This Page Is Inserted by IFW Operations and is not a part of the Official Record

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

IMAGES ARE BEST AVAILABLE COPY.

As rescanning documents will not correct images, please do not report the images to the Image Problem Mailbox.

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date 21 March 2002 (21.03.2002)

PCT

(10) International Publication Number WO 02/22635 A1

- (51) International Patent Classification7: C07H 21/02, 21/04, A61K 48/00, C12Q 1/68, C12P 19/34, C12N 15/85, 15/86
- (21) International Application Number: PCT/US01/28235
- (22) International Filing Date:

10 September 2001 (10.09.2001)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

09/659,791

11 September 2000 (11.09.2000) US

- (71) Applicant (for all designated States except US): ISIS PHARMACEUTICALS, INC. [US/US]; 2292 l²araday Avenue, Carisbad, CA 92008 (US).
- (72) Inventors; and
- (75) Inventors/Applicants (for US only): MONIA, Brett, P. [US/US]; 7605 Nueva Castilla Way, La Costa, CA 92009 (US). FREIER, Susan, M. [US/US]; 2946 Renault Street, San Diego, CA 92122 (US).
- (74) Agents: LICATA, Jane, Massey et al.; Licata & Tyrrell P.C., 66 E. Main Street, Marlton, NJ 08053 (US).

- (81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

Published:

- with international search report
- before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.



22635 A

(54) Title: ANTISENSE MODULATION OF CLUSTERIN EXPRESSION

(57) Abstract: Antisense compounds, compositions and methods are provided for modulating the expression of clusterin. The compositions comprise antisense compounds, particularly antisense oligonucleotides, targeted to nucleic acids encoding clusterin. Methods of using these compounds for modulation of clusterin expression and for treatment of diseases associated with expression of clusterin are provided.

ANTISENSE MODULATION OF CLUSTERIN EXPRESSION

FIELD OF THE INVENTION

The present invention provides compositions and methods for modulating the expression of clusterin. In particular, this invention relates to compounds, particularly oligonucleotides, specifically hybridizable with nucleic acids encoding clusterin. Such compounds have been shown to modulate the expression of clusterin.

BACKGROUND OF THE INVENTION

Clusterin is an amphipathic glycoprotein that was first isolated from the male reproductive system (Bettuzzi et al., Biochem. J., 1989, 257, 293-296; O'Bryan et al., J. Clin. Invest., 1990, 85, 1477-1486). Subsequently, it has been shown that clusterin is ubiquitously distributed among tissues, having a wide range of biologic properties. Investigators from several disciplines, therefore, have isolated clusterin homologs under more than ten different names reviewed in (Bailey and Griswold, Mol. Cell. Endocrinol., 1999, 151, 17-23; Koch-Brandt and Morgans, Prog. Mol. Subcell. Biol., 1996, 16, 130-149; Meri and Jarva, Vox. Sang., 1998, 74, 291-302; Silkensen et al.,

The clusterin protein consists of two non-identical subunits of 34 kDa and 47 kDa, designated alpha and beta, respectively. Clusterin expression is induced almost exclusively as a result of cellular injury, death, or pathology.

Among its many roles, clusterin is a component of the soluble SCb-5 complement complex which is assembled in the plasma upon activation of the complement cascade (Choi et al., Mol. Immunol., 1989, 26, 835-840; Kirszbaum et al., 35 Embo J., 1989, 8, 711-718; Murphy et al., Int. Immunol.,

1989, 1, 551-554; Tschopp and French, Clin. Exp. Immunol.,

—1994, 97 Suppl-2, 11-14). Binding of clusterin has been shown to abolish the membranolytic potential of complement complexes and it has therefore been termed complement lysis inhibitor (CLI) (Jenne and Tschopp, Proc. Natl. Acad. Sci. U. S. A., 1989, 86, 7123-7127).

Further investigations of clusterin demonstrated that it circulates in plasma as a high density lipoprotein (HDL) complex which serves not only as an inhibitor of the lytic 10 complement cascade, but as a regulator of lipid transport and local lipid redistribution (Jenne et al., J. Biol. Chem., 1991, 266, 11030-11036). In this capacity, clusterin isolated and characterized by de Silva et al. and was given the name Apolipoprotein J (ApoJ) (de Silva et 15 al., Biochemistry, 1990, 29, 5380-5389; de Silva et al., J. Biol. Chem., 1990, 265, 13240-13247; de Silva et al., J. Biol. Chem., 1990, 265, 14292-14297). In these studies, clusterin (ApoJ) was shown to play a role in cholesterol transport in the liver and in the regulation of vascular 20 smooth muscle cell differentiation (de Silva et al., J. Biol. Chem., 1990, 265, 13240-13247; Moulson and Millis, J. Cell. Physiol., 1999, 180, 355-364). A link between the modulation of HDL and complement activity is provided by studies by James et al. that characterize the association 25 of a high density lipoprotein, NA1/NA2, with apolipoprotein A-I (ApoA-I). This novel protein NA1/NA2, was subsequently shown to be clusterin (James et al., Arterioscler. Thromb., 1991, 11, 645-652).

Clusterin has also been shown to participate in the cellular process of programmed cell death or apoptosis.

Clusterin expression demarcates cells undergoing apoptosis (Buttyan et al., Mol. Cell. Biol., 1989, 9, 3473-3481) and in studies of the kidney, the onset of hydronephrosis following unilateral obstruction is associated with the

30

increased expression of proteins encoded by the clusterin gene_(Connor et al., Kidney_Int., 1991, 39, 1098-1103)...In both of these studies, clusterin is referred to by two other synonyms, sulfated glycoprotein-2 gene (SGP-2) and 5 testosterone-repressed prostate message-2 (TRPM-2) (Buttyan et al., Mol. Cell. Biol., 1989, 9, 3473-3481; Connor et al., Kidney Int., 1991, 39, 1098-1103).

Sensibar et al. showed that cell death in the prostate, induced by tumor necrosis factor alpha, could be 10 prevented by overexpressing clusterin. In these studies, transfection of LNCaP cells with any of four 21-mer antisense phosphorothioate oligonucleotides targeting the clusterin coding region resulted in an increase of cell death (Sensibar et al., Cancer Res., 1995, 55, 2431-2437).

Miyake et al. further demonstrated the role of 15 clusterin as an anti-apoptotic gene in the Shionogi tumor model, a model used for the study of castration-induced apoptosis (Miyake et al., Cancer Res., 2000, 60, 170-176).

In this model, androgen-dependent mammary carcinoma

20 xenograft tumors in male mice undergo regression after castration but recur as apoptosis-induced tumors after one month. Using a phosphorothicate 21-mer antisense oligonucleotide to the mouse clusterin gene targeting the translation initiation site, Miyake et al. were able to

25 show that treatment with the clusterin antisense oligonucleotide of mice with Shionogi tumors resulted in a more rapid onset of apoptosis and time to complete regresssion. There was also a significant delay of emergence of androgen-independent recurrent tumors compared to control oligonucleotide treated controls.

Using the same oligonucleotide in an experiment designed to test the efficacy of the oligonucleotide in combination with paclitaxel, Miyake et al. showed that the combination of antisense oligonucleotide and paclitaxel 35 induced apoptosis in Shionogi tumors better than either

agent alone. These studies suggest that the antisense oligonucleotide may be useful in enhancing the effects of cytotoxic chemotherapy in hormone-refractory prostate cancer (Miyake et al., Cancer Res., 2000, 60, 2547-2554).

Ten antisense oligodeoxynucleotides targeted to human TRPM-2 (clusterin) were designed by Miyake et al. (Clin. Cancer Res., 2000, 6, 1655-1663) to identify potent oligonucleotides that specifically inhibit TRPM-2 expression in human androgen-independent prostate cancer

10 PC-2 cells. Seven of the ten oligonucleotides had little or no effect on TRPM-2 mRNA expression. The other three oligonucleotides were described by the authors as having moderate effects. The most active oligonucleotide was also tested for ability to enhance the response of PC-3 cells to Taxol or mitoxantrone.

Another antisense oligonucleotide, targeting the AUG initiation codon of clusterin was used to investigate the role of clusterin in endothelial cell activation. In these studies, it was shown that clusterin expression is upregulated upon laminar shear stress and that reduction of clusterin levels via antisense treatment increased endothelial cell activation (Urbich et al., Circulation, 2000, 101, 352-355).

The level of clusterin is increased in the hippocampus and frontal cortex of the brains of Alzheimer's disease patients. It is currently believed that clusterin, by binding to beta-amyloid, a protein known to aggregate in the brains of these patients, acts to link the progression of this disease to the complement system (Choi-Miura and Oda, Neurobiol. Aging, 1996, 17, 717-722).

Most recently, clusterin has been isolated as a KU70 binding protein. KU binding proteins (KUBs) are involved in DNA repair pathways. Clusterin (KUB1) was identified as an autoantigen in serum of patients with scleroderma-

35 polymyositis syndrome and shown to dimerize with KUP80 to

30

form an ATP dependent helicase and a regulatory component of_a_DNA_dependent_protein_kinase_(PRKDC)_involved_in____ double-strand break repair and V(D)J recombination (Yang et al., Nucleic Acids Res., 1999, 27, 2165-2174).

Clusterin is overexpressed in many disease states including neurodegenerative disorders, gliomas, retinitis pigmentosa and expression is induced in acute and chronic models of renal injury and disease, following ureter obstruction, ischemia/reperfusion, and atherosclerosis 10 reviewed in (Silkensen et al., Biochem. Cell. Biol., 1994, The pharmacological modulation of clusterin 72, 483-488). activity and/or expression may therefore be an appropriate point of therapeutic intervention in pathological conditions.

The expression of clusterin, or variants thereof, has 15 been used as a means of differentiating normal versus abnormal cells in the study of male infertility. A method of assessing acrosomal status of sperm morphology comprising contacting a sperm sample with an 20 immunologically reactive molecule which binds to one form

Currently, there are no known therapeutic agents which effectively inhibit the synthesis of clusterin and to date, investigative strategies aimed at modulating clusterin 25 function have involved the use of antibodies, antisense

of clusterin and not another is disclosed in WO 95/16916.

oligonucleotides and chemical inhibitors. There remains, however, a long felt need for additional agents capable of effectively inhibiting

clusterin function. Antisense technology is emerging as an effective means

for reducing the expression of specific gene products and may therefore prove to be uniquely useful in a number of therapeutic, diagnostic, and research applications for the modulation of clusterin expression.

The present invention provides compositions and methods-for modulating-clusterin-expression,-including modulation of the alpha and/or beta subunits.

5 SUMMARY OF THE INVENTION

The present invention is directed to compounds,
particularly antisense oligonucleotides, which are targeted
to a nucleic acid encoding clusterin, and which modulate
the expression of clusterin. Pharmaceutical and other

compositions comprising the compounds of the invention are
also provided. Further provided are methods of modulating
the expression of clusterin in cells or tissues comprising
contacting said cells or tissues with one or more of the
antisense compounds or compositions of the invention.

Further provided are methods of treating an animal,
particularly a human, suspected of having or being prone to
a disease or condition associated with expression of
clusterin by administering a therapeutically or
prophylactically effective amount of one or more of the
antisense compounds or compositions of the invention.

DETAILED DESCRIPTION OF THE INVENTION

The present invention employs oligomeric compounds, particularly antisense oligonucleotides, for use in

25 modulating the function of nucleic acid molecules encoding clusterin, ultimately modulating the amount of clusterin produced. This is accomplished by providing antisense compounds which specifically hybridize with one or more nucleic acids encoding clusterin. As used herein, the

30 terms "target nucleic acid" and "nucleic acid encoding clusterin" encompass DNA encoding clusterin, RNA (including pre-mRNA and mRNA) transcribed from such DNA, and also cDNA derived from such RNA. The specific hybridization of an oligomeric compound with its target nucleic acid interferes with the normal function of the nucleic acid. This

-7-

modulation of function of a target nucleic acid by compounds which specifically hybridize to it is generally_____ referred to as "antisense". The functions of DNA to be interfered with include replication and transcription. The 5 functions of RNA to be interfered with include all vital functions such as, for example, translocation of the RNA to the site of protein translation, translation of protein from the RNA, splicing of the RNA to yield one or more mRNA species, and catalytic activity which may be engaged in or 10 facilitated by the RNA. The overall effect of such interference with target nucleic acid function is modulation of the expression of clusterin. In the context of the present invention, "modulation" means either an increase (stimulation) or a decrease (inhibition) in the 15 expression of a gene. In the context of the present invention, inhibition is the preferred form of modulation of gene expression and mRNA is a preferred target.

It is preferred to target specific nucleic acids for antisense. "Targeting" an antisense compound to a 20 particular nucleic acid, in the context of this invention, is a multistep process. The process usually begins with the identification of a nucleic acid sequence whose function is to be modulated. This may be, for example, a cellular gene (or mRNA transcribed from the gene) whose 25 expression is associated with a particular disorder or disease state, or a nucleic acid molecule from an infectious agent. In the present invention, the target is a nucleic acid molecule encoding clusterin. The targeting process also includes determination of a site or sites 30 within this gene for the antisense interaction to occur such that the desired effect, e.g., detection or modulation of expression of the protein, will result. Within the context of the present invention, a preferred intragenic site is the region encompassing the translation initiation 35 or termination codon of the open reading frame (ORF) of the

gene. Since, as is known in the art, the translation initiation codon is typically 5'-AUG (in transcribed mRNA molecules; 5'-ATG in the corresponding DNA molecule), the translation initiation codon is also referred to as the 5 "AUG codon," the "start codon" or the "AUG start codon". Α minority of genes have a translation initiation codon having the RNA sequence 5'-GUG, 5'-UUG or 5'-CUG, and 5'-AUA, 5'-ACG and 5'-CUG have been shown to function in vivo. Thus, the terms "translation initiation codon" and "start codon" can encompass many codon sequences, even 10 though the initiator amino acid in each instance is typically methionine (in eukaryotes) or formylmethionine (in prokaryotes). It is also known in the art that eukaryotic and prokaryotic genes may have two or more 15 alternative start codons, any one of which may be preferentially utilized for translation initiation in a particular cell type or tissue, or under a particular set of conditions. In the context of the invention, "start codon" and "translation initiation codon" refer to the 20 codon or codons that are used in vivo to initiate translation of an mRNA molecule transcribed from a gene encoding clusterin, regardless of the sequence(s) of such codons.

It is also known in the art that a translation

25 termination codon (or "stop codon") of a gene may have one of three sequences, i.e., 5'-UAA, 5'-UAG and 5'-UGA (the corresponding DNA sequences are 5'-TAA, 5'-TAG and 5'-TGA, respectively). The terms "start codon region" and "translation initiation codon region" refer to a portion of such an mRNA or gene that encompasses from about 25 to about 50 contiguous nucleotides in either direction (i.e., 5' or 3') from a translation initiation codon. Similarly, the terms "stop codon region" and "translation termination codon region" refer to a portion of such an mRNA or gene

35 that encompasses from about 25 to about 50 contiguous

nucleotides in either direction (i.e., 5' or 3') from a translation termination codon.

The open reading frame (ORF) or "coding region," which is known in the art to refer to the region between the 5 translation initiation codon and the translation termination codon, is also a region which may be targeted effectively. Other target regions include the 5' untranslated region (5'UTR), known in the art to refer to the portion of an mRNA in the 5' direction from the 10 translation initiation codon, and thus including nucleotides between the 5' cap site and the translation initiation codon of an mRNA or corresponding nucleotides on the gene, and the 3' untranslated region (3'UTR), known in the art to refer to the portion of an mRNA in the 3' 15 direction from the translation termination codon, and thus including nucleotides between the translation termination codon and 3' end of an mRNA or corresponding nucleotides on the gene. The 5' cap of an mRNA comprises an N7-methylated guanosine residue joined to the 5'-most residue of the mRNA 20 via a 5'-5' triphosphate linkage. The 5' cap region of an mRNA is considered to include the 5' cap structure itself as well as the first 50 nucleotides adjacent to the cap. The 5' cap region may also be a preferred target region.

Although some eukaryotic mRNA transcripts are directly
translated, many contain one or more regions, known as
"introns," which are excised from a transcript before it is
translated. The remaining (and therefore translated)
regions are known as "exons" and are spliced together to
form a continuous mRNA sequence. mRNA splice sites, i.e.,
intron-exon junctions, may also be preferred target
regions, and are particularly useful in situations where
aberrant splicing is implicated in disease, or where an
overproduction of a particular mRNA splice product is
implicated in disease. Aberrant fusion junctions due to
rearrangements or deletions are also preferred targets. It

has also been found that introns can also be effective, and therefore preferred, target regions for antisense compounds targeted, for example, to DNA or pre-mRNA.

-10-

Once one or more target sites have been identified,

oligonucleotides are chosen which are sufficiently
complementary to the target, i.e., hybridize sufficiently
well and with sufficient specificity, to give the desired
effect.

In the context of this invention, "hybridization"

means hydrogen bonding, which may be Watson-Crick,

Hoogsteen or reversed Hoogsteen hydrogen bonding, between

complementary nucleoside or nucleotide bases. For example,

adenine and thymine are complementary nucleobases which

pair through the formation of hydrogen bonds.

- "Complementary," as used herein, refers to the capacity for precise pairing between two nucleotides. For example, if a nucleotide at a certain position of an oligonucleotide is capable of hydrogen bonding with a nucleotide at the same position of a DNA or RNA molecule, then the oligonucleotide and the DNA or RNA are considered to be complementary to each other at that position. The oligonucleotide and the DNA or RNA are complementary to each other when a
- molecule are occupied by nucleotides which can hydrogen

 25 bond with each other. Thus, "specifically hybridizable"

 and "complementary" are terms which are used to indicate a

 sufficient degree of complementarity or precise pairing

 such that stable and specific binding occurs between the

 oligonucleotide and the DNA or RNA target. It is

sufficient number of corresponding positions in each

30 understood in the art that the sequence of an antisense compound need not be 100% complementary to that of its target nucleic acid to be specifically hybridizable. An antisense compound is specifically hybridizable when binding of the compound to the target DNA or RNA molecule interferes with the normal function of the target DNA or

-11-

RNA to cause a loss of utility, and there is a sufficient

degree of complementarity to avoid non-specific binding of
the antisense compound to non-target sequences under
conditions in which specific binding is desired, i.e.,

under physiological conditions in the case of in vivo
assays or therapeutic treatment, and in the case of in
vitro assays, under conditions in which the assays are
performed.

Antisense and other compounds of the invention which

10 hybridize to the target and inhibit expression of the

target are identified through experimentation, and the

sequences of these compounds are hereinbelow identified as

preferred embodiments of the invention. The target sites to

which these preferred sequences are complementary are

15 hereinbelow referred to as "active sites" and are therefore

preferred sites for targeting. Therefore another embodiment

of the invention encompasses compounds which hybridize to

these active sites.

Antisense compounds are commonly used as research

reagents and diagnostics. For example, antisense
oligonucleotides, which are able to inhibit gene expression
with exquisite specificity, are often used by those of
ordinary skill to elucidate the function of particular
genes. Antisense compounds are also used, for example, to
distinguish between functions of various members of a
biological pathway. Antisense modulation has, therefore,
been harnessed for research use.

The specificity and sensitivity of antisense is also harnessed by those of skill in the art for therapeutic

30 uses. Antisense oligonucleotides have been employed as therapeutic moieties in the treatment of disease states in animals and man. Antisense oligonucleotide drugs, including ribozymes, have been safely and effectively administered to humans and numerous clinical trials are presently underway. It is thus established that

WO 02/22635

oligonucleotides can be useful therapeutic modalities that can be configured to be useful in treatment regimes for treatment of cells, tissues and animals, especially humans.

In the context of this invention, the term

5 "oligonucleotide" refers to an oligomer or polymer of ribonucleic acid (RNA) or deoxyribonucleic acid (DNA) or mimetics thereof. This term includes oligonucleotides composed of naturally-occurring nucleobases, sugars and covalent internucleoside (backbone) linkages as well as

10 oligonucleotides having non-naturally-occurring portions which function similarly. Such modified or substituted oligonucleotides are often preferred over native forms because of desirable properties such as, for example, enhanced cellular uptake, enhanced affinity for nucleic acid target and increased stability in the presence of nucleases.

While antisense oligonucleotides are a preferred form of antisense compound, the present invention comprehends other oligomeric antisense compounds, including but not limited to oligonucleotide mimetics such as are described below. The antisense compounds in accordance with this invention preferably comprise from about 8 to about 50 nucleobases (i.e. from about 8 to about 50 linked nucleosides). Particularly preferred antisense compounds are antisense oligonucleotides, even more preferably those comprising from about 12 to about 30 nucleobases. Antisense compounds include ribozymes, external guide sequence (EGS) oligonucleotides (oligozymes), and other short catalytic RNAs or catalytic oligonucleotides which hybridize to the target nucleic acid and modulate its expression.

As is known in the art, a nucleoside is a base-sugar combination. The base portion of the nucleoside is normally a heterocyclic base. The two most common classes of such heterocyclic bases are the purines and the pyrimidines. Nucleotides are nucleosides that further

include a phosphate group covalently linked to the sugar portion of the nucleoside. For those nucleosides that include a pentofuranosyl sugar, the phosphate group can be linked to either the 2', 3' or 5' hydroxyl moiety of the sugar. In forming oligonucleotides, the phosphate groups covalently link adjacent nucleosides to one another to form a linear polymeric compound. In turn the respective ends of this linear polymeric structure can be further joined to form a circular structure, however, open linear structures are generally preferred. Within the oligonucleotide structure, the phosphate groups are commonly referred to as forming the internucleoside backbone of the oligonucleotide. The normal linkage or backbone of RNA and DNA is a 3' to 5' phosphodiester linkage.

useful in this invention include oligonucleotides containing modified backbones or non-natural internucleoside linkages. As defined in this specification, oligonucleotides having modified backbones include those that retain a phosphorus atom in the backbone and those that do not have a phosphorus atom in the backbone. For the purposes of this specification, and as sometimes referenced in the art, modified oligonucleotides that do not have a phosphorus atom in their internucleoside backbone can also be considered to be oligonucleosides.

Preferred modified oligonucleotide backbones include, for example, phosphorothioates, chiral phosphorothioates, phosphorodithioates, phosphotriesters, aminoalkyl-phosphotriesters, methyl and other alkyl phosphonates including 3'-alkylene phosphonates and chiral phosphonates, phosphinates, phosphoramidates including 3'-amino phosphoramidate and aminoalkylphosphoramidates, thionophosphoramidates, thionoalkylphosphorates, thionoalkylphosphotriesters, and boranophosphates having normal 3'-5' linkages, 2'-5' linked analogs of these, and

those having inverted polarity wherein the adjacent pairs of nucleoside units are linked 3'-5' to 5'-3' or 2'-5' to 5'-2'. Various salts, mixed salts and free acid forms are also included.

Representative United States patents that teach the preparation of the above phosphorus-containing linkages include, but are not limited to, U.S.: 3,687,808; 4,469,863; 4,476,301; 5,023,243; 5,177,196; 5,188,897; 5,264,423; 5,276,019; 5,278,302; 5,286,717; 5,321,131; 5,399,676; 5,405,939; 5,453,496; 5,455,233; 5,466,677; 5,476,925; 5,519,126; 5,536,821; 5,541,306; 5,550,111; 5,563,253; 5,571,799; 5,587,361; and 5,625,050, certain of which are commonly owned with this application, and each of which is herein incorporated by reference.

15 Preferred modified oligonucleotide backbones that do not include a phosphorus atom therein have backbones that are formed by short chain alkyl or cycloalkyl internucleoside linkages, mixed heteroatom and alkyl or cycloalkyl internucleoside linkages, or one or more short 20 chain heteroatomic or heterocyclic internucleoside These include those having morpholino linkages linkages. (formed in part from the sugar portion of a nucleoside); siloxane backbones; sulfide, sulfoxide and sulfone backbones; formacetyl and thioformacetyl backbones; 25 methylene formacetyl and thioformacetyl backbones; alkene containing backbones; sulfamate backbones; methyleneimino and methylenehydrazino backbones; sulfonate and sulfonamide backbones; amide backbones; and others having mixed N, O, S and CH2 component parts.

Representative United States patents that teach the preparation of the above oligonucleosides include, but are not limited to, U.S.: 5,034,506; 5,166,315; 5,185,444; 5,214,134; 5,216,141; 5,235,033; 5,264,562; 5,264,564; 5,405,938; 5,434,257; 5,466,677; 5,470,967; 5,489,677; 5,541,307; 5,561,225; 5,596,086; 5,602,240; 5,610,289;

5,602,240; 5,608,046; 5,610,289; 5,618,704; 5,623,070; 5,663,312; 5,633,360; 5,677,437; and 5,677,439, certain of which are commonly owned with this application, and each of which is herein incorporated by reference.

In other preferred oligonucleotide mimetics, both the sugar and the internucleoside linkage, i.e., the backbone, of the nucleotide units are replaced with novel groups. The base units are maintained for hybridization with an appropriate nucleic acid target compound. One such 10 oligomeric compound, an oligonucleotide mimetic that has been shown to have excellent hybridization properties, is referred to as a peptide nucleic acid (PNA). compounds, the sugar-backbone of an oligonucleotide is replaced with an amide containing backbone, in particular 15 an aminoethylglycine backbone. The nucleobases are retained and are bound directly or indirectly to aza nitrogen atoms of the amide portion of the backbone. Representative United States patents that teach the preparation of PNA compounds include, but are not limited 20 to, U.S.: 5,539,082; 5,714,331; and 5,719,262, each of which is herein incorporated by reference. Further teaching of PNA compounds can be found in Nielsen et al., Science, 1991, 254, 1497-1500.

Most preferred embodiments of the invention are oligonucleotides with phosphorothioate backbones and oligonucleosides with heteroatom backbones, and in particular -CH2-NH-O-CH2-, -CH2-N(CH3)-O-CH2- [known as a methylene (methylimino) or MMI backbone], -CH2-O-N(CH3)-CH2-, -CH2-N(CH3)-N(CH3)-CH2- and -O-N(CH3)-CH2-CH2- [wherein the native phosphodiester backbone is represented as -O-P-O-CH2-] of the above referenced U.S. patent 5,489,677, and the amide backbones of the above referenced U.S. patent 5,602,240. Also preferred are oligonucleotides having morpholino backbone structures of the above-referenced U.S. patent 5,034,506.

Modified oligonucleotides may also contain one or more substituted sugar moieties. Preferred oligonucleotides comprise one of the following at the 2' position: OH; F; O-, S-, or N-alkyl; O-, S-, or N-alkenyl; O-, S- or N-5 alkynyl; or O-alkyl-O-alkyl, wherein the alkyl, alkenyl and alkynyl may be substituted or unsubstituted C_1 to C_{10} alkyl or C2 to C10 alkenyl and alkynyl. Particularly preferred are $O[(CH_2)_nO]_mCH_3$, $O(CH_2)_nOCH_3$, $O(CH_2)_nNH_2$, $O(CH_2)_nCH_3$, $O(CH_2)_nONH_2$, and $O(CH_2)_nON[(CH_2)_nCH_3)]_2$, where n and m are 10 from 1 to about 10. Other preferred oligonucleotides comprise one of the following at the 2' position: C1 to C10 lower alkyl, substituted lower alkyl, alkaryl, aralkyl, Oalkaryl or O-aralkyl, SH, SCH3, OCN, Cl, Br, CN, CF3, OCF3, SOCH₃, SO₂CH₃, ONO₂, NO₂, N₃, NH₂, heterocycloalkyl, 15 heterocycloalkaryl, aminoalkylamino, polyalkylamino, substituted silyl, an RNA cleaving group, a reporter group, an intercalator, a group for improving the pharmacokinetic properties of an oligonucleotide, or a group for improving the pharmacodynamic properties of an oligonucleotide, and 20 other substituents having similar properties. A preferred modification includes 2'-methoxyethoxy (2'-O-CH2CH2OCH3, also known as 2'-0-(2-methoxyethyl) or 2'-MOE) (Martin et al., Helv. Chim. Acta, 1995, 78, 486-504) i.e., an alkoxyalkoxy group. A further preferred modification 25 includes 2'-dimethylaminooxyethoxy, i.e., a O(CH₂)₂ON(CH₃)₂ group, also known as 2'-DMAOE, as described in examples hereinbelow, and 2'-dimethylaminoethoxyethoxy (also known in the art as 2'-O-dimethylaminoethoxyethyl or 2'-DMAEOE), i.e., 2'-O-CH2-O-CH2-N(CH2)2, also described in examples 30 hereinbelow.

Other preferred modifications include 2'-methoxy (2'-O-CH₃), 2'-aminopropoxy (2'-OCH₂CH₂CH₂CH₂NH₂) and 2'-fluoro (2'-F). Similar modifications may also be made at other positions on the oligonucleotide, particularly the 3' position of the sugar on the 3' terminal nucleotide or in

2'-5' linked oligonucleotides and the 5' position of 5'
terminal nucleotide. Oligonucleotides may also have sugar
mimetics such as cyclobutyl moieties in place of the
pentofuranosyl sugar. Representative United States patents

5 that teach the preparation of such modified sugar
structures include, but are not limited to, U.S.:
4,981,957; 5,118,800; 5,319,080; 5,359,044; 5,393,878;
5,446,137; 5,466,786; 5,514,785; 5,519,134; 5,567,811;
5,576,427; 5,591,722; 5,597,909; 5,610,300; 5,627,053;
10 5,639,873; 5,646,265; 5,658,873; 5,670,633; and 5,700,920,
certain of which are commonly owned with the instant
application, and each of which is herein incorporated by
reference in its entirety.

Oligonucleotides may also include nucleobase (often referred to in the art simply as "base") modifications or 15 substitutions. As used herein, "unmodified" or "natural" nucleobases include the purine bases adenine (A) and quanine (G), and the pyrimidine bases thymine (T), cytosine (C) and uracil (U). Modified nucleobases include other synthetic and natural nucleobases such as 5-methylcytosine 20 (5-me-C), 5-hydroxymethyl cytosine, xanthine, hypoxanthine, 2-aminoadenine, 6-methyl and other alkyl derivatives of adenine and quanine, 2-propyl and other alkyl derivatives of adenine and guanine, 2-thiouracil, 2-thiothymine and 2-25 thiocytosine, 5-halouracil and cytosine, 5-propynyl uracil and cytosine, 6-azo uracil, cytosine and thymine, 5-uracil (pseudouracil), 4-thiouracil, 8-halo, 8-amino, 8-thiol, 8thioalkyl, 8-hydroxyl and other 8-substituted adenines and guanines, 5-halo particularly 5-bromo, 5-trifluoromethyl 30 and other 5-substituted uracils and cytosines, 7methylquanine and 7-methyladenine, 8-azaguanine and 8azaadenine, 7-deazaguanine and 7-deazaadenine and 3deazaquanine and 3-deazaadenine. Further nucleobases include those disclosed in United States Patent No. 3,687,808, those disclosed in The Concise Encyclopedia Of

Polymer Science And Engineering, pages 858-859, Kroschwitz, J.I., ed. John Wiley & Sons, 1990, those disclosed by Englisch et al., Angewandte Chemie, International Edition, 1991, 30, 613, and those disclosed by Sanghvi, Y.S.,

- 5 Chapter 15, Antisense Research and Applications, pages 289-302, Crooke, S.T. and Lebleu, B., ed., CRC Press, 1993.

 Certain of these nucleobases are particularly useful for increasing the binding affinity of the oligomeric compounds of the invention. These include 5-substituted pyrimidines,
- 6-azapyrimidines and N-2, N-6 and O-6 substituted purines, including 2-aminopropyladenine, 5-propynyluracil and 5-propynylcytosine. 5-methylcytosine substitutions have been shown to increase nucleic acid duplex stability by 0.6-1.2°C (Sanghvi, Y.S., Crooke, S.T. and Lebleu, B., eds.,
- 15 Antisense Research and Applications, CRC Press, Boca Raton, 1993, pp. 276-278) and are presently preferred base substitutions, even more particularly when combined with 2'-0-methoxyethyl sugar modifications.

Representative United States patents that teach the preparation of certain of the above noted modified nucleobases as well as other modified nucleobases include, but are not limited to, the above noted U.S. 3,687,808, as well as U.S.: 4,845,205; 5,130,302; 5,134,066; 5,175,273; 5,367,066; 5,432,272; 5,457,187; 5,459,255; 5,484,908;

5,502,177; 5,525,711; 5,552,540; 5,587,469; 5,594,121, 5,596,091; 5,614,617; and 5,681,941, certain of which are commonly owned with the instant application, and each of which is herein incorporated by reference, and United States patent 5,750,692, which is commonly owned with the instant application and also herein incorporated by reference.

Another modification of the oligonucleotides of the invention involves chemically linking to the oligonucleotide one or more moieties or conjugates which

WO 02/22635

enhance the activity, cellular distribution or cellular uptake of the oligonucleotide. Such moieties include but are not limited to lipid moieties such as a cholesterol moiety (Letsinger et al., Proc. Natl. Acad. Sci. USA, 1989, 5 86, 6553-6556), cholic acid (Manoharan et al., Bioorg. Med. Chem. Let., 1994, 4, 1053-1060), a thioether, e.g., hexyl-S-tritylthiol (Manoharan et al., Ann. N.Y. Acad. Sci., 1992, 660, 306-309; Manoharan et al., Bioorg. Med. Chem. Let., 1993, 3, 2765-2770), a thiocholesterol (Oberhauser et 10 al., Nucl. Acids Res., 1992, 20, 533-538), an aliphatic chain, e.g., dodecandiol or undecyl residues (Saison-Behmoaras et al., EMBO J., 1991, 10, 1111-1118; Kabanov et al., FEBS Lett., 1990, 259, 327-330; Svinarchuk et al., Biochimie, 1993, 75, 49-54), a phospholipid, e.g., di-15 hexadecyl-rac-glycerol or triethylammonium 1,2-di-0hexadecyl-rac-glycero-3-H-phosphonate (Manoharan et al., Tetrahedron Lett., 1995, 36, 3651-3654; Shea et al., Nucl. Acids Res., 1990, 18, 3777-3783), a polyamine or a polyethylene glycol chain (Manoharan et al., Nucleosides & 20 Nucleotides, 1995, 14, 969-973), or adamantane acetic acid (Manoharan et al., Tetrahedron Lett., 1995, 36, 3651-3654), a palmityl moiety (Mishra et al., Biochim. Biophys. Acta, 1995, 1264, 229-237), or an octadecylamine or hexylaminocarbonyl-oxycholesterol moiety (Crooke et al., J.

25 Pharmacol. Exp. Ther., 1996, 277, 923-937.

Representative United States patents that teach the preparation of such oligonucleotide conjugates include, but are not limited to, U.S.: 4,828,979; 4,948,882; 5,218,105; 5,525,465; 5,541,313; 5,545,730; 5,552,538; 5,578,717, 30 5,580,731; 5,580,731; 5,591,584; 5,109,124; 5,118,802; 5,138,045; 5,414,077; 5,486,603; 5,512,439; 5,578,718; 5,608,046; 4,587,044; 4,605,735; 4,667,025; 4,762,779; 4,789,737; 4,824,941; 4,835,263; 4,876,335; 4,904,582; 4,958,013; 5,082,830; 5,112,963; 5,214,136; 5,082,830;

5,112,963; 5,214,136; 5,245,022; 5,254,469; 5,258,506;
5,262,536; 5,272,250; 5,292,873; 5,317,098; 5,371,241,
5,391,723; 5,416,203, 5,451,463; 5,510,475; 5,512,667;
5,514,785; 5,565,552; 5,567,810; 5,574,142; 5,585,481;
5,587,371; 5,595,726; 5,597,696; 5,599,923; 5,599,928 and
5,688,941, certain of which are commonly owned with the instant application, and each of which is herein incorporated by reference.

It is not necessary for all positions in a given

compound to be uniformly modified, and in fact more than
one of the aforementioned modifications may be incorporated
in a single compound or even at a single nucleoside within
an oligonucleotide. The present invention also includes
antisense compounds which are chimeric compounds.

"Chimeric" antisense compounds or "chimeras," in the

- 15 "Chimeric" antisense compounds or "chimeras," in the context of this invention, are antisense compounds, particularly oligonucleotides, which contain two or more chemically distinct regions, each made up of at least one monomer unit, i.e., a nucleotide in the case of an
- oligonucleotide compound. These oligonucleotides typically contain at least one region wherein the oligonucleotide is modified so as to confer upon the oligonucleotide increased resistance to nuclease degradation, increased cellular uptake, and/or increased binding affinity for the target
- nucleic acid. An additional region of the oligonucleotide may serve as a substrate for enzymes capable of cleaving RNA:DNA or RNA:RNA hybrids. By way of example, RNase H is a cellular endonuclease which cleaves the RNA strand of an RNA:DNA duplex. Activation of RNase H, therefore, results
- in cleavage of the RNA target, thereby greatly enhancing the efficiency of oligonucleotide inhibition of gene expression. Consequently, comparable results can often be obtained with shorter oligonucleotides when chimeric oligonucleotides are used, compared to phosphorothicate
- 35 deoxyoligonucleotides hybridizing to the same target

region. Cleavage of the RNA target can be routinely detected by gel electrophoresis and, if necessary, associated nucleic acid hybridization techniques known in the art.

Chimeric antisense compounds of the invention may be formed as composite structures of two or more oligonucleotides, modified oligonucleotides, oligonucleosides and/or oligonucleotide mimetics as described above. Such compounds have also been referred to in the art as hybrids or gapmers. Representative United States patents that teach the preparation of such hybrid structures include, but are not limited to, U.S.: 5,013,830; 5,149,797; 5,220,007; 5,256,775; 5,366,878; 5,403,711; 5,491,133; 5,565,350; 5,623,065; 5,652,355; 15 5,652,356; and 5,700,922, certain of which are commonly

owned with the instant application, and each of which is

herein incorporated by reference in its entirety.

The antisense compounds used in accordance with this invention may be conveniently and routinely made through

the well-known technique of solid phase synthesis.

Equipment for such synthesis is sold by several vendors including, for example, Applied Biosystems (Foster City, CA). Any other means for such synthesis known in the art may additionally or alternatively be employed. It is well known to use similar techniques to prepare oligonucleotides

The antisense compounds of the invention are synthesized in vitro and do not include antisense compositions of biological origin, or genetic vector constructs designed to direct the in vivo synthesis of antisense molecules.

such as the phosphorothioates and alkylated derivatives.

The compounds of the invention may also be admixed, encapsulated, conjugated or otherwise associated with other molecules, molecule structures or mixtures of compounds, as for example, liposomes, receptor targeted molecules, oral,

rectal, topical or other formulations, for assisting in uptake, distribution and/or absorption. Representative United States patents that teach the preparation of such uptake, distribution and/or absorption assisting

5 formulations include, but are not limited to, U.S.:
5,108,921; 5,354,844; 5,416,016; 5,459,127; 5,521,291;
5,543,158; 5,547,932; 5,583,020; 5,591,721; 4,426,330;
4,534,899; 5,013,556; 5,108,921; 5,213,804; 5,227,170;
5,264,221; 5,356,633; 5,395,619; 5,416,016; 5,417,978;
10 5,462,854; 5,469,854; 5,512,295; 5,527,528; 5,534,259;
5,543,152; 5,556,948; 5,580,575; and 5,595,756, each of which is herein incorporated by reference.

The antisense compounds of the invention encompass any pharmaceutically acceptable salts, esters, or salts of such esters, or any other compound which, upon administration to an animal including a human, is capable of providing (directly or indirectly) the biologically active metabolite or residue thereof. Accordingly, for example, the disclosure is also drawn to prodrugs and pharmaceutically acceptable salts of the compounds of the invention, pharmaceutically acceptable salts of such prodrugs, and other bioequivalents.

The term "prodrug" indicates a therapeutic agent that is prepared in an inactive form that is converted to an active form (i.e., drug) within the body or cells thereof by the action of endogenous enzymes or other chemicals and/or conditions. In particular, prodrug versions of the oligonucleotides of the invention are prepared as SATE [(S-acetyl-2-thioethyl) phosphate] derivatives according to the methods disclosed in WO 93/24510 to Gosselin et al., published December 9, 1993 or in WO 94/26764 and U.S. 5,770,713 to Imbach et al.

The term "pharmaceutically acceptable salts" refers to physiologically and pharmaceutically acceptable salts of the compounds of the invention: i.e., salts that retain the

WO 02/22635

desired biological activity of the parent compound and do

not impart undesired toxicological effects thereto.

Pharmaceutically acceptable base addition salts are formed with metals or amines, such as alkali and alkaline 5 earth metals or organic amines. Examples of metals used as cations are sodium, potassium, magnesium, calcium, and the like. Examples of suitable amines are N, N'-dibenzylethylenediamine, chloroprocaine, choline, diethanolamine, dicyclohexylamine, ethylenediamine, 10 N-methylglucamine, and procaine (see, for example, Berge et al., "Pharmaceutical Salts," J. of Pharma Sci., 1977, 66, 1-19). The base addition salts of said acidic compounds are prepared by contacting the free acid form with a sufficient amount of the desired base to produce the salt 15 in the conventional manner. The free acid form may be regenerated by contacting the salt form with an acid and isolating the free acid in the conventional manner. free acid forms differ from their respective salt forms somewhat in certain physical properties such as solubility 20 in polar solvents, but otherwise the salts are equivalent to their respective free acid for purposes of the present invention. As used herein, a "pharmaceutical addition salt" includes a pharmaceutically acceptable salt of an acid form of one of the components of the compositions of 25 the invention. These include organic or inorganic acid salts of the amines. Preferred acid salts are the hydrochlorides, acetates, salicylates, nitrates and phosphates. Other suitable pharmaceutically acceptable salts are well known to those skilled in the art and 30 include basic salts of a variety of inorganic and organic acids, such as, for example, with inorganic acids, such as for example hydrochloric acid, hydrobromic acid, sulfuric acid or phosphoric acid; with organic carboxylic, sulfonic, sulfo or phospho acids or N-substituted sulfamic acids, for 35 example acetic acid, propionic acid, glycolic acid,

succinic acid, maleic acid, hydroxymaleic acid, methylmaleic acid, fumaric acid, malic acid, tartaric acid, lactic acid, oxalic acid, gluconic acid, glucaric acid, glucuronic acid, citric acid, benzoic acid, cinnamic acid, 5 mandelic acid, salicylic acid, 4-aminosalicylic acid, 2-phenoxybenzoic acid, 2-acetoxybenzoic acid, embonic acid, nicotinic acid or isonicotinic acid; and with amino acids, such as the 20 alpha-amino acids involved in the synthesis of proteins in nature, for example glutamic acid or 10 aspartic acid, and also with phenylacetic acid, methanesulfonic acid, ethanesulfonic acid, 2-hydroxyethanesulfonic acid, ethane-1,2-disulfonic acid, benzenesulfonic acid, 4-methylbenzenesulfonic acid, naphthalene-2-sulfonic acid, naphthalene-1,5-disulfonic 15 acid, 2- or 3-phosphoglycerate, glucose-6-phosphate, N-cyclohexylsulfamic acid (with the formation of cyclamates), or with other acid organic compounds, such as ascorbic acid. Pharmaceutically acceptable salts of compounds may also be prepared with a pharmaceutically 20 acceptable cation. Suitable pharmaceutically acceptable cations are well known to those skilled in the art and include alkaline, alkaline earth, ammonium and quaternary ammonium cations. Carbonates or hydrogen carbonates are also possible.

25 For oligonucleotides, preferred examples of pharmaceutically acceptable salts include but are not limited to (a) salts formed with cations such as sodium, potassium, ammonium, magnesium, calcium, polyamines such as spermine and spermidine, etc.; (b) acid addition salts 30 formed with inorganic acids, for example hydrochloric acid, hydrobromic acid, sulfuric acid, phosphoric acid, nitric acid and the like; (c) salts formed with organic acids such as, for example, acetic acid, oxalic acid, tartaric acid, succinic acid, maleic acid, fumaric acid, gluconic acid, citric acid, malic acid, ascorbic acid, benzoic acid,

WO 02/22635

tannic acid, palmitic acid, alginic acid, polyglutamic

acid, naphthalenesulfonic acid, methanesulfonic acid,
p-toluenesulfonic acid, naphthalenedisulfonic acid,
polygalacturonic acid, and the like; and (d) salts formed

from elemental anions such as chlorine, bromine, and
iodine.

-25-

The antisense compounds of the present invention can be utilized for diagnostics, therapeutics, prophylaxis and as research reagents and kits. For therapeutics, an 10 animal, preferably a human, suspected of having a disease or disorder which can be treated by modulating the expression of clusterin is treated by administering antisense compounds in accordance with this invention. The compounds of the invention can be utilized in 15 pharmaceutical compositions by adding an effective amount of an antisense compound to a suitable pharmaceutically acceptable diluent or carrier. Use of the antisense compounds and methods of the invention may also be useful prophylactically, e.g., to prevent or delay infection, inflammation or tumor formation, for example. 20

The antisense compounds of the invention are useful for research and diagnostics, because these compounds hybridize to nucleic acids encoding clusterin, enabling sandwich and other assays to easily be constructed to exploit this fact. Hybridization of the antisense oligonucleotides of the invention with a nucleic acid encoding clusterin can be detected by means known in the art. Such means may include conjugation of an enzyme to the oligonucleotide, radiolabelling of the oligonucleotide or any other suitable detection means. Kits using such detection means for detecting the level of clusterin in a sample may also be prepared.

The present invention also includes pharmaceutical compositions and formulations which include the antisense compounds of the invention. The pharmaceutical

-26-

compositions of the present invention may be administered in a number of ways depending upon whether local or systemic treatment is desired and upon the area to be treated. Administration may be topical (including ophthalmic and to mucous membranes including vaginal and rectal delivery), pulmonary, e.g., by inhalation or insufflation of powders or aerosols, including by nebulizer; intratracheal, intranasal, epidermal and transdermal), oral or parenteral. Parenteral administration includes intravenous, intraarterial, subcutaneous, intraperitoneal or intramuscular injection or infusion; or intracranial, e.g., intrathecal or intraventricular, administration. Oligonucleotides with at least one 2'-O-methoxyethyl modification are believed to be particularly useful for oral administration.

Pharmaceutical compositions and formulations for topical administration may include transdermal patches, ointments, lotions, creams, gels, drops, suppositories, sprays, liquids and powders. Conventional pharmaceutical carriers, aqueous, powder or oily bases, thickeners and the like may be necessary or desirable. Coated condoms, gloves and the like may also be useful.

Compositions and formulations for oral administration include powders or granules, suspensions or solutions in water or non-aqueous media, capsules, sachets or tablets. Thickeners, flavoring agents, diluents, emulsifiers, dispersing aids or binders may be desirable.

Compositions and formulations for parenteral, intrathecal or intraventricular administration may include sterile aqueous solutions which may also contain buffers, diluents and other suitable additives such as, but not limited to, penetration enhancers, carrier compounds and other pharmaceutically acceptable carriers or excipients.

Pharmaceutical compositions of the present invention include, but are not limited to, solutions, emulsions, and

-27-

liposome-containing formulations. These compositions may be generated from a variety of components that include, but are not limited to, preformed liquids, self-emulsifying solids and self-emulsifying semisolids.

The pharmaceutical formulations of the present invention, which may conveniently be presented in unit dosage form, may be prepared according to conventional techniques well known in the pharmaceutical industry. Such techniques include the step of bringing into association

10 the active ingredients with the pharmaceutical carrier(s) or excipient(s). In general the formulations are prepared by uniformly and intimately bringing into association the active ingredients with liquid carriers or finely divided solid carriers or both, and then, if necessary, shaping the product.

The compositions of the present invention may be formulated into any of many possible dosage forms such as, but not limited to, tablets, capsules, liquid syrups, soft gels, suppositories, and enemas. The compositions of the present invention may also be formulated as suspensions in aqueous, non-aqueous or mixed media. Aqueous suspensions may further contain substances which increase the viscosity of the suspension including, for example, sodium carboxymethylcellulose, sorbitol and/or dextran. The suspension may also contain stabilizers.

In one embodiment of the present invention the pharmaceutical compositions may be formulated and used as foams. Pharmaceutical foams include formulations such as, but not limited to, emulsions, microemulsions, creams,

30 jellies and liposomes. While basically similar in nature these formulations vary in the components and the consistency of the final product. The preparation of such compositions and formulations is generally known to those skilled in the pharmaceutical and formulation arts and may

-28-

be applied to the formulation of the compositions of the present invention.

Emulsions

- 5 The compositions of the present invention may be prepared and formulated as emulsions. Emulsions are typically heterogenous systems of one liquid dispersed in another in the form of droplets usually exceeding 0.1 μm in diameter. (Idson, in *Pharmaceutical Dosage Forms*,
- 10 Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 199; Rosoff, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., Volume 1, p. 245; Block in Pharmaceutical Dosage Forms, Lieberman,
- 15 Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 2, p. 335; Higuchi et al., in Remington's Pharmaceutical Sciences, Mack Publishing Co., Easton, PA, 1985, p. 301). Emulsions are often biphasic systems comprising of two immiscible liquid phases
- intimately mixed and dispersed with each other. In general, emulsions may be either water-in-oil (w/o) or of the oil-in-water (o/w) variety. When an aqueous phase is finely divided into and dispersed as minute droplets into a bulk oily phase the resulting composition is called a
- water-in-oil (w/o) emulsion. Alternatively, when an oily phase is finely divided into and dispersed as minute droplets into a bulk aqueous phase the resulting composition is called an oil-in-water (o/w) emulsion. Emulsions may contain additional components in addition to
- the dispersed phases and the active drug which may be present as a solution in either the aqueous phase, oily phase or itself as a separate phase. Pharmaceutical excipients such as emulsifiers, stabilizers, dyes, and anti-oxidants may also be present in emulsions as needed.

Pharmaceutical emulsions may also be multiple emulsions
that are comprised of more than two phases such as, for
example, in the case of oil-in-water-in-oil (o/w/o) and
water-in-oil-in-water (w/o/w) emulsions. Such complex
formulations often provide certain advantages that simple
binary emulsions do not. Multiple emulsions in which
individual oil droplets of an o/w emulsion enclose small
water droplets constitute a w/o/w emulsion. Likewise a
system of oil droplets enclosed in globules of water
stabilized in an oily continuous provides an o/w/o
emulsion.

Emulsions are characterized by little or no thermodynamic stability. Often, the dispersed or discontinuous phase of the emulsion is well dispersed into 15 the external or continuous phase and maintained in this form through the means of emulsifiers or the viscosity of the formulation. Either of the phases of the emulsion may be a semisolid or a solid, as is the case of emulsion-style ointment bases and creams. Other means of stabilizing 20 emulsions entail the use of emulsifiers that may be incorporated into either phase of the emulsion. Emulsifiers may broadly be classified into four categories: synthetic surfactants, naturally occurring emulsifiers, absorption bases, and finely dispersed solids (Idson, in 25 Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 199).

Synthetic surfactants, also known as surface active agents, have found wide applicability in the formulation of emulsions and have been reviewed in the literature (Rieger, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 285; Idson, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), Marcel Dekker, Inc., New York, N.Y., 1988, volume 1, p. 199). Surfactants are

-30-

typically amphiphilic and comprise a hydrophilic and a hydrophobic portion. The ratio of the hydrophilic to the hydrophobic nature of the surfactant has been termed the hydrophile/lipophile balance (HLB) and is a valuable tool in categorizing and selecting surfactants in the preparation of formulations. Surfactants may be classified into different classes based on the nature of the hydrophilic group: nonionic, anionic, cationic and amphoteric (Rieger, in *Pharmaceutical Dosage Forms*,

10 Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 285).

Naturally occurring emulsifiers used in emulsion formulations include lanolin, beeswax, phosphatides, lecithin and acacia. Absorption bases possess hydrophilic properties such that they can soak up water to form w/o emulsions yet retain their semisolid consistencies, such as anhydrous lanolin and hydrophilic petrolatum. Finely divided solids have also been used as good emulsifiers especially in combination with surfactants and in viscous preparations. These include polar inorganic solids, such as heavy metal hydroxides, nonswelling clays such as bentonite, attapulgite, hectorite, kaolin, montmorillonite, colloidal aluminum silicate and colloidal magnesium aluminum silicate, pigments and nonpolar solids such as carbon or glyceryl tristearate.

A large variety of non-emulsifying materials are also included in emulsion formulations and contribute to the properties of emulsions. These include fats, oils, waxes, fatty acids, fatty alcohols, fatty esters, humectants, hydrophilic colloids, preservatives and antioxidants (Block, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 335; Idson, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 199).

WO 02/22635

PCT/US01/28235

Hydrophilic colloids or hydrocolloids include

naturally occurring gums and synthetic polymers such as
polysaccharides (for example, acacia, agar, alginic acid,
carrageenan, guar gum, karaya gum, and tragacanth),

5 cellulose derivatives (for example, carboxymethylcellulose
and carboxypropylcellulose), and synthetic polymers (for
example, carbomers, cellulose ethers, and carboxyvinyl
polymers). These disperse or swell in water to form
colloidal solutions that stabilize emulsions by forming

10 strong interfacial films around the dispersed-phase
droplets and by increasing the viscosity of the external
phase.

Since emulsions often contain a number of ingredients such as carbohydrates, proteins, sterols and phosphatides

that may readily support the growth of microbes, these formulations often incorporate preservatives. Commonly used preservatives included in emulsion formulations include methyl paraben, propyl paraben, quaternary ammonium salts, benzalkonium chloride, esters of p-hydroxybenzoic

acid, and boric acid. Antioxidants are also commonly added to emulsion formulations to prevent deterioration of the formulation. Antioxidants used may be free radical scavengers such as tocopherols, alkyl gallates, butylated hydroxyanisole, butylated hydroxytoluene, or reducing

agents such as ascorbic acid and sodium metabisulfite, and antioxidant synergists such as citric acid, tartaric acid, and lecithin.

The application of emulsion formulations via dermatological, oral and parenteral routes and methods for their manufacture have been reviewed in the literature (Idson, in *Pharmaceutical Dosage Forms*, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 199). Emulsion formulations for oral delivery have been very widely used because of reasons of ease of formulation, efficacy from an absorption and

35

bioavailability standpoint. (Rosoff, in Pharmaceutical Dosage-Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 245; Idson, in Pharmaceutical Dosage Forms, Lieberman, Rieger 5 and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 199). Mineral-oil base laxatives, oilsoluble vitamins and high fat nutritive preparations are among the materials that have commonly been administered orally as o/w emulsions.

10 In one embodiment of the present invention, the compositions of oligonucleotides and nucleic acids are formulated as microemulsions. A microemulsion may be defined as a system of water, oil and amphiphile which is a single optically isotropic and thermodynamically stable liquid solution (Rosoff, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 245). Typically microemulsions are systems that are prepared by first dispersing an oil in an aqueous surfactant solution and 20 then adding a sufficient amount of a fourth component, generally an intermediate chain-length alcohol to form a transparent system. Therefore, microemulsions have also been described as thermodynamically stable, isotropically clear dispersions of two immiscible liquids that are stabilized by interfacial films of surface-active molecules (Leung and Shah, in: Controlled Release of Drugs: Polymers and Aggregate Systems, Rosoff, M., Ed., 1989, VCH Publishers, New York, pages 185-215). Microemulsions commonly are prepared via a combination of three to five components that include oil, water, surfactant, cosurfactant and electrolyte. Whether the microemulsion is of the water-in-oil (w/o) or an oil-in-water (o/w) type is dependent on the properties of the oil and surfactant used and on the structure and geometric packing of the polar heads and hydrocarbon tails of the surfactant molecules

(Schott, in Remington's Pharmaceutical Sciences, Mack
Publishing Co., Easton, PA, 1985, p. 271).

The phenomenological approach utilizing phase diagrams has been extensively studied and has yielded a

5 comprehensive knowledge, to one skilled in the art, of how to formulate microemulsions (Rosoff, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 245; Block, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 335). Compared to conventional emulsions, microemulsions offer the advantage of solubilizing water-insoluble drugs in a formulation of thermodynamically stable droplets that are formed spontaneously.

Surfactants used in the preparation of microemulsions include, but are not limited to, ionic surfactants, nonionic surfactants, Brij 96, polyoxyethylene oleyl ethers, polyglycerol fatty acid esters, tetraglycerol monolaurate (ML310), tetraglycerol monooleate (MO310), hexaglycerol 20 monooleate (PO310), hexaglycerol pentaoleate (PO500), decaglycerol monocaprate (MCA750), decaglycerol monooleate (MO750), decaglycerol sequioleate (SO750), decaglycerol decaoleate (DAO750), alone or in combination with 25 cosurfactants. The cosurfactant, usually a short-chain alcohol such as ethanol, 1-propanol, and 1-butanol, serves to increase the interfacial fluidity by penetrating into the surfactant film and consequently creating a disordered film because of the void space generated among surfactant 30 molecules. Microemulsions may, however, be prepared without the use of cosurfactants and alcohol-free selfemulsifying microemulsion systems are known in the art. The aqueous phase may typically be, but is not limited to, water, an aqueous solution of the drug, glycerol, PEG300, 35 PEG400, polyglycerols, propylene glycols, and derivatives

PCT/US01/28235

of ethylene glycol. The oil phase may include, but is not limited to, materials such as Captex 300, Captex 355, Capmul MCM, fatty acid esters, medium chain (C8-C12) mono, di, and tri-glycerides, polyoxyethylated glyceryl fatty acid esters, fatty alcohols, polyglycolized glycerides, saturated polyglycolized C8-C10 glycerides, vegetable oils and silicone oil.

-34-

Microemulsions are particularly of interest from the standpoint of drug solubilization and the enhanced absorption of drugs. Lipid based microemulsions (both o/w and w/o) have been proposed to enhance the oral bioavailability of drugs, including peptides (Constantinides et al., Pharmaceutical Research, 1994, 11, 1385-1390; Ritschel, Meth. Find. Exp. Clin. Pharmacol., 1993, 13, 205). Microemulsions afford advantages of improved drug solubilization, protection of drug from enzymatic hydrolysis, possible enhancement of drug absorption due to surfactant-induced alterations in membrane fluidity and permeability, ease of preparation, 20 ease of oral administration over solid dosage forms, improved clinical potency, and decreased toxicity (Constantinides et al., Pharmaceutical Research, 1994, 11, 1385; Ho et al., J. Pharm. Sci., 1996, 85, 138-143). Often microemulsions may form spontaneously when their components 25 are brought together at ambient temperature. This may be particularly advantageous when formulating thermolabile drugs, peptides or oligonucleotides. Microemulsions have also been effective in the transdermal delivery of active components in both cosmetic and pharmaceutical applications. It is expected that the microemulsion compositions and formulations of the present invention will facilitate the increased systemic absorption of

oligonucleotides and nucleic acids from the

gastrointestinal tract, as well as improve the local cellular uptake of oligonucleotides and nucleic acids

within the gastrointestinal tract, vagina, buccal cavity and other areas of administration.

Microemulsions of the present invention may also contain additional components and additives such as

5 sorbitan monostearate (Grill 3), Labrasol, and penetration enhancers to improve the properties of the formulation and to enhance the absorption of the oligonucleotides and nucleic acids of the present invention. Penetration enhancers used in the microemulsions of the present

10 invention may be classified as belonging to one of five broad categories - surfactants, fatty acids, bile salts, chelating agents, and non-chelating non-surfactants (Lee et al., Critical Reviews in Therapeutic Drug Carrier Systems, 1991, p. 92). Each of these classes has been discussed above.

Liposomes

There are many organized surfactant structures besides microemulsions that have been studied and used for the formulation of drugs. These include monolayers, micelles, bilayers and vesicles. Vesicles, such as liposomes, have attracted great interest because of their specificity and the duration of action they offer from the standpoint of drug delivery. As used in the present invention, the term "liposome" means a vesicle composed of amphiphilic lipids arranged in a spherical bilayer or bilayers.

Liposomes are unilamellar or multilamellar vesicles which have a membrane formed from a lipophilic material and an aqueous interior. The aqueous portion contains the composition to be delivered. Cationic liposomes possess the advantage of being able to fuse to the cell wall. Non-cationic liposomes, although not able to fuse as efficiently with the cell wall, are taken up by macrophages in vivo.

In order to cross intact mammalian skin, lipid

vesicles must pass through a series of fine pores, each

with a diameter less than 50 nm, under the influence of a

suitable transdermal gradient. Therefore, it is desirable

to use a liposome which is highly deformable and able to

pass through such fine pores.

Further advantages of liposomes include; liposomes obtained from natural phospholipids are biocompatible and biodegradable; liposomes can incorporate a wide range of water and lipid soluble drugs; liposomes can protect encapsulated drugs in their internal compartments from metabolism and degradation (Rosoff, in Pharmaceutical Dosage Forms, Lieberman, Rieger and Banker (Eds.), 1988, Marcel Dekker, Inc., New York, N.Y., volume 1, p. 245).

15 Important considerations in the preparation of liposome formulations are the lipid surface charge, vesicle size and the aqueous volume of the liposomes.

Liposomes are useful for the transfer and delivery of active ingredients to the site of action. Because the liposomal membrane is structurally similar to biological membranes, when liposomes are applied to a tissue, the liposomes start to merge with the cellular membranes. As the merging of the liposome and cell progresses, the liposomal contents are emptied into the cell where the active agent may act.

Liposomal formulations have been the focus of extensive investigation as the mode of delivery for many drugs. There is growing evidence that for topical administration, liposomes present several advantages over other formulations. Such advantages include reduced side-effects related to high systemic absorption of the administered drug, increased accumulation of the administered drug at the desired target, and the ability to administer a wide variety of drugs, both hydrophilic and hydrophobic, into the skin.

30

Several reports have detailed the ability of liposomes to deliver agents including high-molecular weight DNA into the skin. Compounds including analgesics, antibodies, hormones and high-molecular weight DNAs have been 5 administered to the skin. The majority of applications resulted in the targeting of the upper epidermis.

Liposomes fall into two broad classes. Cationic liposomes are positively charged liposomes which interact with the negatively charged DNA molecules to form a stable 10 complex. The positively charged DNA/liposome complex binds to the negatively charged cell surface and is internalized in an endosome. Due to the acidic pH within the endosome, the liposomes are ruptured, releasing their contents into the cell cytoplasm (Wang et al., Biochem. Biophys. Res. 15 Commun., 1987, 147, 980-985).

Liposomes which are pH-sensitive or negatively-charged, entrap DNA rather than complex with it. Since both the DNA and the lipid are similarly charged, repulsion rather than complex formation occurs.

20 Nevertheless, some DNA is entrapped within the aqueous interior of these liposomes. pH-sensitive liposomes have been used to deliver DNA encoding the thymidine kinase gene to cell monolayers in culture. Expression of the exogenous gene was detected in the target cells (Zhou et al., Journal 25 of Controlled Release, 1992, 19, 269-274).

One major type of liposomal composition includes phospholipids other than naturally-derived phosphatidylcholine. Neutral liposome compositions, for example, can be formed from dimyristoyl phosphatidylcholine (DMPC) or dipalmitoyl phosphatidylcholine (DPPC). Anionic liposome compositions generally are formed from dimyristoyl phosphatidylglycerol, while anionic fusogenic liposomes are formed primarily from dioleoyl phosphatidylethanolamine (DOPE). Another type of liposomal composition is formed 35 from phosphatidylcholine (PC) such as, for example, soybean PC, and egg PC. Another type is formed from mixtures of phospholipid and/or phosphatidylcholine and/or cholesterol.

Several studies have assessed the topical delivery of liposomal drug formulations to the skin. Application of liposomes containing interferon to guinea pig skin resulted in a reduction of skin herpes sores while delivery of interferon via other means (e.g. as a solution or as an emulsion) were ineffective (Weiner et al., Journal of Drug Targeting, 1992, 2, 405-410). Further, an additional study tested the efficacy of interferon administered as part of a liposomal formulation to the administration of interferon using an aqueous system, and concluded that the liposomal formulation was superior to aqueous administration (du Plessis et al., Antiviral Research, 1992, 18, 259-265).

Non-ionic liposomal systems have also been examined to determine their utility in the delivery of drugs to the skin, in particular systems comprising non-ionic surfactant and cholesterol. Non-ionic liposomal formulations comprising NovasomeTM I (glyceryl

dilaurate/cholesterol/polyoxyethylene-10-stearyl ether) and
NovasomeTM II (glyceryl distearate/
 cholesterol/polyoxyethylene-10-stearyl ether) were used to
 deliver cyclosporin-A into the dermis of mouse skin.
 Results indicated that such non-ionic liposomal systems
were effective in facilitating the deposition of
 cyclosporin-A into different layers of the skin (Hu et al.
 S.T.P.Pharma. Sci., 1994, 4, 6, 466).

Liposomes also include "sterically stabilized"
liposomes, a term which, as used herein, refers to
liposomes comprising one or more specialized lipids that,
when incorporated into liposomes, result in enhanced
circulation lifetimes relative to liposomes lacking such
specialized lipids. Examples of sterically stabilized
liposomes are those in which part of the vesicle-forming

(Lim et al.).

lipid portion of the liposome (A) comprises one or more glycolipids, such as monosialoganglioside G_{M1}, or (B) is derivatized with one or more hydrophilic polymers, such as a polyethylene glycol (PEG) moiety. While not wishing to 5 be bound by any particular theory, it is thought in the art that, at least for sterically stabilized liposomes containing gangliosides, sphingomyelin, or PEG-derivatized lipids, the enhanced circulation half-life of these sterically stabilized liposomes derives from a reduced 10 uptake into cells of the reticuloendothelial system (RES) (Allen et al., FEBS Letters, 1987, 223, 42; Wu et al., Cancer Research, 1993, 53, 3765). Various liposomes comprising one or more glycolipids are known in the art. Papahadjopoulos et al. (Ann. N.Y. Acad. Sci., 1987, 507, 15 64) reported the ability of monosialoganglioside G_{M1} , galactocerebroside sulfate and phosphatidylinositol to improve blood half-lives of liposomes. These findings were expounded upon by Gabizon et al. (Proc. Natl. Acad. Sci. U.S.A., 1988, 85, 6949). U.S. Patent No. 4,837,028 and WO 20 88/04924, both to Allen et al., disclose liposomes comprising (1) sphingomyelin and (2) the ganglioside G_{M1} or a galactocerebroside sulfate ester. U.S. Patent No. 5,543,152 (Webb et al.) discloses liposomes comprising sphingomyelin. Liposomes comprising 1,2-sn-25 dimyristoylphosphatidylcholine are disclosed in WO 97/13499

Many liposomes comprising lipids derivatized with one or more hydrophilic polymers, and methods of preparation thereof, are known in the art. Sunamoto et al. (Bull.

30 Chem. Soc. Jpn., 1980, 53, 2778) described liposomes comprising a nonionic detergent, 2C₁₂15G, that contains a PEG moiety. Illum et al. (FEBS Lett., 1984, 167, 79) noted that hydrophilic coating of polystyrene particles with polymeric glycols results in significantly enhanced blood

half-lives. Synthetic phospholipids modified by the attachment of carboxylic groups of polyalkylene glycols (e.g., PEG) are described by Sears (U.S. Patent Nos. 4,426,330 and 4,534,899). Klibanov et al. (FEBS Lett., 5 1990, 268, 235) described experiments demonstrating that liposomes comprising phosphatidylethanolamine (PE) derivatized with PEG or PEG stearate have significant increases in blood circulation half-lives. Blume et al. (Biochimica et Biophysica Acta, 1990, 1029, 91) extended 10 such observations to other PEG-derivatized phospholipids, e.g., DSPE-PEG, formed from the combination of distearoylphosphatidylethanolamine (DSPE) and PEG. Liposomes having covalently bound PEG moieties on their external surface are described in European Patent No. EP 0 15 445 131 B1 and WO 90/04384 to Fisher. Liposome compositions containing 1-20 mole percent of PE derivatized with PEG, and methods of use thereof, are described by Woodle et al. (U.S. Patent Nos. 5,013,556 and 5,356,633) and Martin et al. (U.S. Patent No. 5,213,804 and European 20 Patent No. EP 0 496 813 B1). Liposomes comprising a number of other lipid-polymer conjugates are disclosed in WO 91/05545 and U.S. Patent No. 5,225,212 (both to Martin et al.) and in WO 94/20073 (Zalipsky et al.) Liposomes comprising PEG-modified ceramide lipids are described in WO 25 96/10391 (Choi et al.). U.S. Patent Nos. 5,540,935 (Miyazaki et al.) and 5,556,948 (Tagawa et al.) describe PEG-containing liposomes that can be further derivatized with functional moieties on their surfaces.

A limited number of liposomes comprising nucleic acids

30 are known in the art. WO 96/40062 to Thierry et al.

discloses methods for encapsulating high molecular weight
nucleic acids in liposomes. U.S. Patent No. 5,264,221 to
Tagawa et al. discloses protein-bonded liposomes and
asserts that the contents of such liposomes may include an

35 antisense RNA. U.S. Patent No. 5,665,710 to Rahman et al.

30

describes certain methods of encapsulating oligodeoxynucleotides in liposomes. WO 97/04787 to Love et al. discloses liposomes comprising antisense oligonucleotides targeted to the raf gene.

Transfersomes are yet another type of liposomes, and 5 are highly deformable lipid aggregates which are attractive candidates for drug delivery vehicles. Transfersomes may be described as lipid droplets which are so highly deformable that they are easily able to penetrate through 10 pores which are smaller than the droplet. Transfersomes are adaptable to the environment in which they are used, e.g. they are self-optimizing (adaptive to the shape of pores in the skin), self-repairing, frequently reach their targets without fragmenting, and often self-loading. 15 make transfersomes it is possible to add surface edgeactivators, usually surfactants, to a standard liposomal composition. Transfersomes have been used to deliver serum albumin to the skin. The transfersome-mediated delivery of serum albumin has been shown to be as effective as 20 subcutaneous injection of a solution containing serum albumin.

Surfactants find wide application in formulations such as emulsions (including microemulsions) and liposomes. most common way of classifying and ranking the properties 25 of the many different types of surfactants, both natural and synthetic, is by the use of the hydrophile/lipophile balance (HLB). The nature of the hydrophilic group (also known as the "head") provides the most useful means for categorizing the different surfactants used in formulations (Rieger, in Pharmaceutical Dosage Forms, Marcel Dekker, Inc., New York, NY, 1988, p. 285).

If the surfactant molecule is not ionized, it is classified as a nonionic surfactant. Nonionic surfactants find wide application in pharmaceutical and cosmetic 35 products and are usable over a wide range of pH values. Ιn general their HLB values range from 2 to about 18 depending on their structure. Nonionic surfactants include nonionic esters such as ethylene glycol esters, propylene glycol esters, glyceryl esters, polyglyceryl esters, sorbitan esters, sucrose esters, and ethoxylated esters. Nonionic alkanolamides and ethers such as fatty alcohol ethoxylates, propoxylated alcohols, and ethoxylated/propoxylated block polymers are also included in this class. The polyoxyethylene surfactants are the most popular members of the nonionic surfactant class.

If the surfactant molecule carries a negative charge when it is dissolved or dispersed in water, the surfactant is classified as anionic. Anionic surfactants include carboxylates such as soaps, acyl lactylates, acyl amides of amino acids, esters of sulfuric acid such as alkyl sulfates and ethoxylated alkyl sulfates, sulfonates such as alkyl benzene sulfonates, acyl isethionates, acyl taurates and sulfosuccinates, and phosphates. The most important members of the anionic surfactant class are the alkyl sulfates and the soaps.

If the surfactant molecule carries a positive charge when it is dissolved or dispersed in water, the surfactant is classified as cationic. Cationic surfactants include quaternary ammonium salts and ethoxylated amines. The quaternary ammonium salts are the most used members of this class.

If the surfactant molecule has the ability to carry either a positive or negative charge, the surfactant is classified as amphoteric. Amphoteric surfactants include acrylic acid derivatives, substituted alkylamides, Nalkylbetaines and phosphatides.

The use of surfactants in drug products, formulations and in emulsions has been reviewed (Rieger, in
Pharmaceutical Dosage Forms, Marcel Dekker, Inc., New York,
NY, 1988, p. 285).

Penetration Enhancers

In one embodiment, the present invention employs various penetration enhancers to effect the efficient

5 delivery of nucleic acids, particularly oligonucleotides, to the skin of animals. Most drugs are present in solution in both ionized and nonionized forms. However, usually only lipid soluble or lipophilic drugs readily cross cell membranes. It has been discovered that even non-lipophilic drugs may cross cell membranes if the membrane to be crossed is treated with a penetration enhancer. In addition to aiding the diffusion of non-lipophilic drugs across cell membranes, penetration enhancers also enhance the permeability of lipophilic drugs.

15 Penetration enhancers may be classified as belonging to one of five broad categories, i.e., surfactants, fatty acids, bile salts, chelating agents, and non-chelating non-surfactants (Lee et al., Critical Reviews in Therapeutic Drug Carrier Systems, 1991, p.92). Each of the above 20 mentioned classes of penetration enhancers are described below in greater detail.

Surfactants: In connection with the present invention, surfactants (or "surface-active agents") are chemical
25 entities which, when dissolved in an aqueous solution, reduce the surface tension of the solution or the interfacial tension between the aqueous solution and another liquid, with the result that absorption of oligonucleotides through the mucosa is enhanced. In
30 addition to bile salts and fatty acids, these penetration enhancers include, for example, sodium lauryl sulfate, polyoxyethylene-9-lauryl ether and polyoxyethylene-20-cetyl ether) (Lee et al., Critical Reviews in Therapeutic Drug Carrier Systems, 1991, p.92); and perfluorochemical

emulsions, such as FC-43. Takahashi et al., J. Pharm.

Pharmacol., 1988, 40, 252).

Fatty acids: Various fatty acids and their derivatives which act as penetration enhancers include, for example, 5 oleic acid, lauric acid, capric acid (n-decanoic acid), myristic acid, palmitic acid, stearic acid, linoleic acid, linolenic acid, dicaprate, tricaprate, monoolein (1monooleoyl-rac-glycerol), dilaurin, caprylic acid, arachidonic acid, glycerol 1-monocaprate, 1-10 dodecylazacycloheptan-2-one, acylcarnitines, acylcholines, C_{1-10} alkyl esters thereof (e.g., methyl, isopropyl and tbutyl), and mono- and di-glycerides thereof (i.e., oleate, laurate, caprate, myristate, palmitate, stearate, linoleate, etc.) (Lee et al., Critical Reviews in 15 Therapeutic Drug Carrier Systems, 1991, p.92; Muranishi, Critical Reviews in Therapeutic Drug Carrier Systems, 1990, 7, 1-33; El Hariri et al., J. Pharm. Pharmacol., 1992, 44, 651-654).

20 Bile salts: The physiological role of bile includes the facilitation of dispersion and absorption of lipids and fat-soluble vitamins (Brunton, Chapter 38 in: Goodman & Gilman's The Pharmacological Basis of Therapeutics, 9th Ed., Hardman et al. Eds., McGraw-Hill, New York, 1996, pp. 25 934-935). Various natural bile salts, and their synthetic derivatives, act as penetration enhancers. Thus the term "bile salts" includes any of the naturally occurring components of bile as well as any of their synthetic derivatives. The bile salts of the invention include, for 30 example, cholic acid (or its pharmaceutically acceptable sodium salt, sodium cholate), dehydrocholic acid (sodium dehydrocholate), deoxycholic acid (sodium deoxycholate), glucholic acid (sodium glucholate), glycholic acid (sodium glycocholate), glycodeoxycholic acid (sodium

glycodeoxycholate), taurocholic acid (sodium taurocholate),
taurodeoxycholic acid (sodium taurodeoxycholate),
chenodeoxycholic acid (sodium chenodeoxycholate),
ursodeoxycholic acid (UDCA), sodium tauro-24,25-dihydro5 fusidate (STDHF), sodium glycodihydrofusidate and
polyoxyethylene-9-lauryl ether (POE) (Lee et al., Critical
Reviews in Therapeutic Drug Carrier Systems, 1991, page 92;
Swinyard, Chapter 39 In: Remington's Pharmaceutical
Sciences, 18th Ed., Gennaro, ed., Mack Publishing Co.,
10 Easton, PA, 1990, pages 782-783; Muranishi, Critical
Reviews in Therapeutic Drug Carrier Systems, 1990, 7, 1-33;
Yamamoto et al., J. Pharm. Exp. Ther., 1992, 263, 25;
Yamashita et al., J. Pharm. Sci., 1990, 79, 579-583).

Chelating Agents: Chelating agents, as used in 15 connection with the present invention, can be defined as compounds that remove metallic ions from solution by forming complexes therewith, with the result that absorption of oligonucleotides through the mucosa is 20 enhanced. With regards to their use as penetration enhancers in the present invention, chelating agents have the added advantage of also serving as DNase inhibitors, as most characterized DNA nucleases require a divalent metal ion for catalysis and are thus inhibited by chelating agents (Jarrett, J. Chromatogr., 1993, 618, 315-339). Chelating agents of the invention include but are not limited to disodium ethylenediaminetetraacetate (EDTA), citric acid, salicylates (e.g., sodium salicylate, 5methoxysalicylate and homovanilate), N-acyl derivatives of 30 collagen, laureth-9 and N-amino acyl derivatives of betadiketones (enamines) (Lee et al., Critical Reviews in Therapeutic Drug Carrier Systems, 1991, page 92; Muranishi, Critical Reviews in Therapeutic Drug Carrier Systems, 1990, 7, 1-33; Buur et al., J. Control Rel., 1990, 14, 43-51).

Non-chelating non-surfactants: As used herein, nonchelating non-surfactant penetration enhancing compounds
can be defined as compounds that demonstrate insignificant
activity as chelating agents or as surfactants but that
nonetheless enhance absorption of oligonucleotides through
the alimentary mucosa (Muranishi, Critical Reviews in
Therapeutic Drug Carrier Systems, 1990, 7, 1-33). This
class of penetration enhancers include, for example,
unsaturated cyclic ureas, 1-alkyl- and 1-alkenylazacycloalkanone derivatives (Lee et al., Critical Reviews in
Therapeutic Drug Carrier Systems, 1991, page 92); and nonsteroidal anti-inflammatory agents such as diclofenac
sodium, indomethacin and phenylbutazone (Yamashita et al.,

J. Pharm. Pharmacol., 1987, 39, 621-626).

Agents that enhance uptake of oligonucleotides at the cellular level may also be added to the pharmaceutical and other compositions of the present invention. For example, cationic lipids, such as lipofectin (Junichi et al, U.S. Patent No. 5,705,188), cationic glycerol derivatives, and

polycationic molecules, such as polylysine (Lollo *et al.*, PCT Application WO 97/30731), are also known to enhance the cellular uptake of oligonucleotides.

Other agents may be utilized to enhance the

25 penetration of the administered nucleic acids, including glycols such as ethylene glycol and propylene glycol, pyrrols such as 2-pyrrol, azones, and terpenes such as limonene and menthone.

30 Carriers

Certain compositions of the present invention also incorporate carrier compounds in the formulation. As used herein, "carrier compound" or "carrier" can refer to a nucleic acid, or analog thereof, which is inert (i.e., does not possess biological activity per se) but is recognized

WO 02/22635

PCT/US01/28235

as a nucleic acid by *in vivo* processes that reduce the bioavailability of a nucleic acid having biological activity by, for example, degrading the biologically active

nucleic acid or promoting its removal from circulation.

5 The coadministration of a nucleic acid and a carrier

The coadministration of a nucleic acid and a carrier compound, typically with an excess of the latter substance, can result in a substantial reduction of the amount of nucleic acid recovered in the liver, kidney or other extracirculatory reservoirs, presumably due to competition

10 between the carrier compound and the nucleic acid for a common receptor. For example, the recovery of a partially phosphorothicate oligonucleotide in hepatic tissue can be reduced when it is coadministered with polyinosinic acid, dextran sulfate, polycytidic acid or 4-acetamido-

15 4'isothiocyano-stilbene-2,2'-disulfonic acid (Miyao et al.,
Antisense Res. Dev., 1995, 5, 115-121; Takakura et al.,
Antisense & Nucl. Acid Drug Dev., 1996, 6, 177-183).

Excipients

In contrast to a carrier compound, a "pharmaceutical carrier" or "excipient" is a pharmaceutically acceptable solvent, suspending agent or any other pharmacologically inert vehicle for delivering one or more nucleic acids to an animal. The excipient may be liquid or solid and is

selected, with the planned manner of administration in mind, so as to provide for the desired bulk, consistency, etc., when combined with a nucleic acid and the other components of a given pharmaceutical composition. Typical pharmaceutical carriers include, but are not limited to,

binding agents (e.g., pregelatinized maize starch,
 polyvinylpyrrolidone or hydroxypropyl methylcellulose,
 etc.); fillers (e.g., lactose and other sugars,
 microcrystalline cellulose, pectin, gelatin, calcium
 sulfate, ethyl cellulose, polyacrylates or calcium hydrogen

35 phosphate, etc.); lubricants (e.g., magnesium stearate,

-48-

talc, silica, colloidal silicon dioxide, stearic acid,
metallic stearates, hydrogenated vegetable oils, corn
starch, polyethylene glycols, sodium benzoate, sodium
acetate, etc.); disintegrants (e.g., starch, sodium starch
glycolate, etc.); and wetting agents (e.g., sodium lauryl
sulphate, etc.).

Pharmaceutically acceptable organic or inorganic excipient suitable for non-parenteral administration which do not deleteriously react with nucleic acids can also be used to formulate the compositions of the present invention. Suitable pharmaceutically acceptable carriers include, but are not limited to, water, salt solutions, alcohols, polyethylene glycols, gelatin, lactose, amylose, magnesium stearate, talc, silicic acid, viscous paraffin, hydroxymethylcellulose, polyvinylpyrrolidone and the like.

Formulations for topical administration of nucleic acids may include sterile and non-sterile aqueous solutions, non-aqueous solutions in common solvents such as alcohols, or solutions of the nucleic acids in liquid or solid oil bases. The solutions may also contain buffers, diluents and other suitable additives. Pharmaceutically acceptable organic or inorganic excipients suitable for non-parenteral administration which do not deleteriously react with nucleic acids can be used.

Suitable pharmaceutically acceptable excipients include, but are not limited to, water, salt solutions, alcohol, polyethylene glycols, gelatin, lactose, amylose, magnesium stearate, talc, silicic acid, viscous paraffin, hydroxymethylcellulose, polyvinylpyrrolidone and the like.

Other Components

25

30

The compositions of the present invention may additionally contain other adjunct components conventionally found in pharmaceutical compositions, at their art-established usage levels. Thus, for example, the

compositions may contain additional, compatible, pharmaceutically-active materials such as, for example, antipruritics, astringents, local anesthetics or anti-inflammatory agents, or may contain additional 5 materials useful in physically formulating various dosage forms of the compositions of the present invention, such as dyes, flavoring agents, preservatives, antioxidants, opacifiers, thickening agents and stabilizers. However, such materials, when added, should not unduly interfere 10 with the biological activities of the components of the compositions of the present invention. The formulations can be sterilized and, if desired, mixed with auxiliary agents, e.g., lubricants, preservatives, stabilizers, wetting agents, emulsifiers, salts for influencing osmotic 15 pressure, buffers, colorings, flavorings and/or aromatic substances and the like which do not deleteriously interact with the nucleic acid(s) of the formulation.

Aqueous suspensions may contain substances which increase the viscosity of the suspension including, for example, sodium carboxymethylcellulose, sorbitol and/or dextran. The suspension may also contain stabilizers.

Certain embodiments of the invention provide
pharmaceutical compositions containing (a) one or more
antisense compounds and (b) one or more other

25 chemotherapeutic agents which function by a non-antisense
mechanism. Examples of such chemotherapeutic agents
include, but are not limited to, anticancer drugs such as
daunorubicin, dactinomycin, doxorubicin, bleomycin,
mitomycin, nitrogen mustard, chlorambucil, melphalan,

30 cyclophosphamide, 6-mercaptopurine, 6-thioguanine,
cytarabine (CA), 5-fluorouracil (5-FU), floxuridine
(5-FUdR), methotrexate (MTX), colchicine, vincristine,
vinblastine, etoposide, teniposide, cisplatin and
diethylstilbestrol (DES). See, generally, The Merck Manual
of Diagnosis and Therapy, 15th Ed., Berkow et al., eds.,

In another related embodiment, compositions of the invention may contain one or more antisense compounds, particularly oligonucleotides, targeted to a first nucleic acid and one or more additional antisense compounds targeted to a second nucleic acid target. Numerous examples of antisense compounds are known in the art. Two or more combined compounds may be used together or sequentially.

The formulation of therapeutic compositions and their 20 subsequent administration is believed to be within the skill of those in the art. Dosing is dependent on severity and responsiveness of the disease state to be treated, with the course of treatment lasting from several days to several months, or until a cure is effected or a diminution 25 of the disease state is achieved. Optimal dosing schedules can be calculated from measurements of drug accumulation in the body of the patient. Persons of ordinary skill can easily determine optimum dosages, dosing methodologies and repetition rates. Optimum dosages may vary depending on 30 the relative potency of individual oligonucleotides, and can generally be estimated based on EC50s found to be effective in in vitro and in vivo animal models. general, dosage is from 0.01 ug to 100 g per kg of body weight, and may be given once or more daily, weekly, 35 monthly or yearly, or even once every 2 to 20 years.

-51-

Persons of ordinary skill in the art can easily estimate

repetition rates for dosing based on measured residence

times and concentrations of the drug in bodily fluids or

tissues. Following successful treatment, it may be

desirable to have the patient undergo maintenance therapy

to prevent the recurrence of the disease state, wherein the

oligonucleotide is administered in maintenance doses,

ranging from 0.01 ug to 100 g per kg of body weight, once

or more daily, to once every 20 years.

While the present invention has been described with specificity in accordance with certain of its preferred embodiments, the following examples serve only to illustrate the invention and are not intended to limit the same.

15

EXAMPLES

Example 1

Nucleoside Phosphoramidites for Oligonucleotide Synthesis Deoxy and 2'-alkoxy amidites

- 2'-Deoxy and 2'-methoxy beta-cyanoethyldiisopropyl phosphoramidites were purchased from commercial sources (e.g. Chemgenes, Needham MA or Glen Research, Inc. Sterling VA). Other 2'-O-alkoxy substituted nucleoside amidites are prepared as described in U.S. Patent 5,506,351, herein
- 25 incorporated by reference. For oligonucleotides synthesized using 2'-alkoxy amidites, the standard cycle for unmodified oligonucleotides was utilized, except the wait step after pulse delivery of tetrazole and base was increased to 360 seconds.
- Oligonucleotides containing 5-methyl-2'-deoxycytidine
 (5-Me-C) nucleotides were synthesized according to
 published methods [Sanghvi, et. al., Nucleic Acids
 Research, 1993, 21, 3197-3203] using commercially available
 phosphoramidites (Glen Research, Sterling VA or ChemGenes,
 Needham MA).

2'-Fluoro amidites

-2'-Fluorodeoxyadenosine amidites

2'-fluoro oligonucleotides were synthesized as described previously [Kawasaki, et. al., J. Med. Chem., 5 1993, 36, 831-841] and United States patent 5,670,633, herein incorporated by reference. Briefly, the protected nucleoside N6-benzoyl-2'-deoxy-2'-fluoroadenosine was synthesized utilizing commercially available 9-beta-Darabinofuranosyladenine as starting material and by 10 modifying literature procedures whereby the 2'-alpha-fluoro atom is introduced by a S_N2-displacement of a 2'-beta-trityl group. Thus N6-benzoyl-9-beta-D-arabinofuranosyladenine was selectively protected in moderate yield as the 3',5'ditetrahydropyranyl (THP) intermediate. Deprotection of 15 the THP and N6-benzoyl groups was accomplished using standard methodologies and standard methods were used to obtain the 5'-dimethoxytrityl-(DMT) and 5'-DMT-3'phosphoramidite intermediates.

20 2'-Fluorodeoxyguanosine

The synthesis of 2'-deoxy-2'-fluoroguanosine was accomplished using tetraisopropyldisiloxanyl (TPDS) protected 9-beta-D-arabinofuranosylguanine as starting material, and conversion to the intermediate diisobutyryl-arabinofuranosylguanosine. Deprotection of the TPDS group was followed by protection of the hydroxyl group with THP to give diisobutyryl di-THP protected arabinofuranosylguanine. Selective O-deacylation and triflation was followed by treatment of the crude product with fluoride, then deprotection of the THP groups. Standard methodologies were used to obtain the 5'-DMT- and 5'-DMT-3'-phosphoramidites.

2'-Fluorouridine

35 Synthesis of 2'-deoxy-2'-fluorouridine was

accomplished by the modification of a literature procedure in which 2,2'-anhydro-1-beta-D-arabinofuranosyluracil was treated with 70% hydrogen fluoride-pyridine. Standard procedures were used to obtain the 5'-DMT and 5'-DMT-5 3'phosphoramidites.

2'-Fluorodeoxycytidine

20

2'-deoxy-2'-fluorocytidine was synthesized via amination of 2'-deoxy-2'-fluorouridine, followed by 10 selective protection to give N4-benzoyl-2'-deoxy-2'fluorocytidine. Standard procedures were used to obtain the 5'-DMT and 5'-DMT-3'phosphoramidites.

2'-0-(2-Methoxyethyl) modified amidites

2'-O-Methoxyethyl-substituted nucleoside amidites are prepared as follows, or alternatively, as per the methods of Martin, P., Helvetica Chimica Acta, 1995, 78, 486-504.

2,2'-Anhydro[1-(beta-D-arabinofuranosyl)-5-methyluridine]

5-Methyluridine (ribosylthymine, commercially available through Yamasa, Choshi, Japan) (72.0 g, 0.279 M), diphenylcarbonate (90.0 g, 0.420 M) and sodium bicarbonate (2.0 g, 0.024 M) were added to DMF (300 mL). The mixture was heated to reflux, with stirring, allowing the evolved carbon dioxide gas to be released in a controlled manner. After 1 hour, the slightly darkened solution was concentrated under reduced pressure. The resulting syrup was poured into diethylether (2.5 L), with stirring. The product formed a gum. The ether was decanted and the residue was dissolved in a minimum amount of methanol (ca. 400 mL). The solution was poured into fresh ether (2.5 L) to yield a stiff gum. The ether was decanted and the gum was dried in a vacuum oven (60°C at 1 mm Hg for 24 h) to

give a solid that was crushed to a light tan powder (57 g, 85% crude yield). The NMR spectrum was consistent with the structure, contaminated with phenol as its sodium salt (ca. 5%). The material was used as is for further reactions (or it can be purified further by column chromatography using a gradient of methanol in ethyl acetate (10-25%) to give a white solid, mp 222-4°C).

2'-O-Methoxyethyl-5-methyluridine

10 2,2'-Anhydro-5-methyluridine (195 g, 0.81 M), tris(2methoxyethyl) borate (231 g, 0.98 M) and 2-methoxyethanol (1.2 L) were added to a 2 L stainless steel pressure vessel and placed in a pre-heated oil bath at 160°C. After heating for 48 hours at 155-160°C, the vessel was opened and the 15 solution evaporated to dryness and triturated with MeOH (200 mL). The residue was suspended in hot acetone (1 L). The insoluble salts were filtered, washed with acetone (150 mL) and the filtrate evaporated. The residue (280 g) was dissolved in CH3CN (600 mL) and evaporated. A silica 20 gel column (3 kg) was packed in CH₂Cl₂/acetone/MeOH (20:5:3) containing 0.5% Et3NH. The residue was dissolved in CH2Cl2 (250 mL) and adsorbed onto silica (150 g) prior to loading onto the column. The product was eluted with the packing solvent to give 160 g (63%) of product. Additional 25 material was obtained by reworking impure fractions.

2'-O-Methoxyethyl-5'-O-dimethoxytrityl-5-methyluridine

2'-O-Methoxyethyl-5-methyluridine (160 g, 0.506 M) was co-evaporated with pyridine (250 mL) and the dried residue dissolved in pyridine (1.3 L). A first aliquot of dimethoxytrityl chloride (94.3 g, 0.278 M) was added and the mixture stirred at room temperature for one hour. A second aliquot of dimethoxytrityl chloride (94.3 g, 0.278 M) was added and the reaction stirred for an additional one

-55-

hour. Methanol (170 mL) was then added to stop the reaction. HPLC showed the presence of approximately 70% product. The solvent was evaporated and triturated with CH₃CN (200 mL). The residue was dissolved in CHCl₃ (1.5 L) and extracted with 2x500 mL of saturated NaHCO₃ and 2x500 mL of saturated NaCl. The organic phase was dried over Na₂SO₄, filtered and evaporated. 275 g of residue was obtained. The residue was purified on a 3.5 kg silica gel column, packed and eluted with EtOAc/hexane/acetone (5:5:1) containing 0.5% Et₃NH. The pure fractions were evaporated to give 164 g of product. Approximately 20 g additional was obtained from the impure fractions to give a total yield of 183 g (57%).

3'-O-Acetyl-2'-O-methoxyethyl-5'-O-dimethoxytrityl-5methyluridine

2'-O-Methoxyethyl-5'-O-dimethoxytrityl-5-methyluridine (106 g, 0.167 M), DMF/pyridine (750 mL of a 3:1 mixture prepared from 562 mL of DMF and 188 mL of pyridine) and 20 acetic anhydride (24.38 mL, 0.258 M) were combined and stirred at room temperature for 24 hours. The reaction was monitored by TLC by first quenching the TLC sample with the addition of MeOH. Upon completion of the reaction, as judged by TLC, MeOH (50 mL) was added and the mixture 25 evaporated at 35°C. The residue was dissolved in CHCl₃ (800 mL) and extracted with 2x200 mL of saturated sodium bicarbonate and 2x200 mL of saturated NaCl. layers were back extracted with 200 mL of CHCl₃. combined organics were dried with sodium sulfate and 30 evaporated to give 122 g of residue (approx. 90% product). The residue was purified on a 3.5 kg silica gel column and eluted using EtOAc/hexane(4:1). Pure product fractions were evaporated to yield 96 g (84%). An additional 1.5 g was recovered from later fractions.

3'-0-Acetyl-2'-O-methoxyethyl-5'-O-dimethoxytrityl-5-methyl-4-triazoleuridine

A first solution was prepared by dissolving 3'-0acetyl-2'-O-methoxyethyl-5'-O-dimethoxytrityl-5-5 methyluridine (96 g, 0.144 M) in CH₃CN (700 mL) and set aside. Triethylamine (189 mL, 1.44 M) was added to a solution of triazole (90 g, 1.3 M) in CH₃CN (1 L), cooled to -5°C and stirred for 0.5 h using an overhead stirrer. POCl₃ was added dropwise, over a 30 minute period, to the stirred 10 solution maintained at 0-10°C, and the resulting mixture stirred for an additional 2 hours. The first solution was added dropwise, over a 45 minute period, to the latter solution. The resulting reaction mixture was stored overnight in a cold room. Salts were filtered from the 15 reaction mixture and the solution was evaporated. The residue was dissolved in EtOAc (1 L) and the insoluble solids were removed by filtration. The filtrate was washed with 1x300 mL of NaHCO3 and 2x300 mL of saturated NaCl, dried over sodium sulfate and evaporated. The residue was 20 triturated with EtOAc to give the title compound.

2'-O-Methoxyethyl-5'-O-dimethoxytrityl-5-methylcytidine

A solution of 3'-O-acetyl-2'-O-methoxyethyl-5'-O
25 dimethoxytrityl-5-methyl-4-triazoleuridine (103 g, 0.141 M)

in dioxane (500 mL) and NH₄OH (30 mL) was stirred at room

temperature for 2 hours. The dioxane solution was

evaporated and the residue azeotroped with MeOH (2x200 mL).

The residue was dissolved in MeOH (300 mL) and transferred

30 to a 2 liter stainless steel pressure vessel. MeOH (400 mL) saturated with NH₃ gas was added and the vessel heated

to 100°C for 2 hours (TLC showed complete conversion). The vessel contents were evaporated to dryness and the residue was dissolved in EtOAc (500 mL) and washed once with

saturated NaCl (200 mL). The organics were dried over sodium sulfate and the solvent was evaporated to give 85 g (95%) of the title compound.

5 N4-Benzoyl-2'-O-methoxyethyl-5'-O-dimethoxytrityl-5-methylcytidine

2'-O-Methoxyethyl-5'-O-dimethoxytrityl-5-methylcytidine (85 g, 0.134 M) was dissolved in DMF (800 mL) and
benzoic anhydride (37.2 g, 0.165 M) was added with

10 stirring. After stirring for 3 hours, TLC showed the
reaction to be approximately 95% complete. The solvent was
evaporated and the residue azeotroped with MeOH (200 mL).
The residue was dissolved in CHCl₃ (700 mL) and extracted
with saturated NaHCO₃ (2x300 mL) and saturated NaCl (2x300

15 mL), dried over MgSO₄ and evaporated to give a residue (96
g). The residue was chromatographed on a 1.5 kg silica
column using EtOAc/hexane (1:1) containing 0.5% Et₃NH as the
eluting solvent. The pure product fractions were
evaporated to give 90 g (90%) of the title compound.

20

N4-Benzoyl-2'-O-methoxyethyl-5'-O-dimethoxytrityl-5-methylcytidine-3'-amidite

M4-Benzoyl-2'-O-methoxyethyl-5'-O-dimethoxytrityl-5-methylcytidine (74 g, 0.10 M) was dissolved in CH₂Cl₂ (1 L).

Tetrazole diisopropylamine (7.1 g) and 2-cyanoethoxytetra(isopropyl)phosphite (40.5 mL, 0.123 M) were added with stirring, under a nitrogen atmosphere. The resulting mixture was stirred for 20 hours at room temperature (TLC showed the reaction to be 95% complete). The reaction

mixture was extracted with saturated NaHCO₃ (1x300 mL) and saturated NaCl (3x300 mL). The aqueous washes were backextracted with CH₂Cl₂ (300 mL), and the extracts were combined, dried over MgSO₄ and concentrated. The residue obtained was chromatographed on a 1.5 kg silica column

using EtOAc/hexane (3:1) as the eluting solvent. The pure

fractions were combined to give 90.6 g (87%) of the title compound.

2'-O-(Aminooxyethyl) nucleoside amidites and 2'-O5 (dimethylaminooxyethyl) nucleoside amidites

2'-(Dimethylaminooxyethoxy) nucleoside amidites

2'-(Dimethylaminooxyethoxy) nucleoside amidites [also known in the art as 2'-0-(dimethylaminooxyethyl) nucleoside amidites] are prepared as described in the following paragraphs. Adenosine, cytidine and guanosine nucleoside amidites are prepared similarly to the thymidine (5-methyluridine) except the exocyclic amines are protected with a benzoyl moiety in the case of adenosine and cytidine and with isobutyryl in the case of guanosine.

5'-O-tert-Butyldiphenylsilyl-O²-2'-anhydro-5-methyluridine

O²-2'-anhydro-5-methyluridine (Pro. Bio. Sint., Varese, 20 Italy, 100.0g, 0.416 mmol), dimethylaminopyridine (0.66g, 0.013eg, 0.0054mmol) were dissolved in dry pyridine (500 ml) at ambient temperature under an argon atmosphere and with mechanical stirring. tert-Butyldiphenylchlorosilane (125.8g, 119.0mL, 1.1eq, 0.458mmol) was added in one 25 portion. The reaction was stirred for 16 h at ambient temperature. TLC (Rf 0.22, ethyl acetate) indicated a complete reaction. The solution was concentrated under reduced pressure to a thick oil. This was partitioned between dichloromethane (1 L) and saturated sodium 30 bicarbonate (2x1 L) and brine (1 L). The organic layer was dried over sodium sulfate and concentrated under reduced pressure to a thick oil. The oil was dissolved in a 1:1 mixture of ethyl acetate and ethyl ether (600mL) and the solution was cooled to

-59~

-10°C. The resulting crystalline product was collected by filtration, washed with ethyl ether (3x200 mL) and dried (40°C, 1mm Hg, 24 h) to 149g (74.8%) of white solid. TLC and NMR were consistent with pure product.

5

5'-O-tert-Butyldiphenylsilyl-2'-O-(2-hydroxyethyl)-5-methyluridine

In a 2 L stainless steel, unstirred pressure reactor was added borane in tetrahydrofuran (1.0 M, 2.0 eq, 622 10 mL). In the fume hood and with manual stirring, ethylene glycol (350 mL, excess) was added cautiously at first until the evolution of hydrogen gas subsided. 5'-O-tert-Butyldiphenylsilyl-02-2'-anhydro-5-methyluridine (149 g, 0.311 mol) and sodium bicarbonate (0.074 g, 0.003 eq) were 15 added with manual stirring. The reactor was sealed and heated in an oil bath until an internal temperature of 160 °C was reached and then maintained for 16 h (pressure < 100 psig). The reaction vessel was cooled to ambient and opened. TLC (Rf 0.67 for desired product and Rf 0.82 for 20 ara-T side product, ethyl acetate) indicated about 70% conversion to the product. In order to avoid additional side product formation, the reaction was stopped, concentrated under reduced pressure (10 to 1mm Hg) in a warm water bath (40-100°C) with the more extreme conditions 25 used to remove the ethylene glycol. [Alternatively, once the low boiling solvent is gone, the remaining solution can be partitioned between ethyl acetate and water. product will be in the organic phase.] The residue was purified by column chromatography (2kg silica gel, ethyl 30 acetate-hexanes gradient 1:1 to 4:1). The appropriate fractions were combined, stripped and dried to product as a white crisp foam (84g, 50%), contaminated starting material (17.4g) and pure reusable starting material 20g. The yield based on starting material less pure recovered starting

material was 58%. TLC and NMR were consistent with 99%

2'-O-([2-phthalimidoxy)ethyl]-5'-t-butyldiphenylsilyl-5-methyluridine

5'-O-tert-Butyldiphenylsilyl-2'-O-(2-hydroxyethyl)-5methyluridine (20g, 36.98mmol) was mixed with triphenylphosphine (11.63g, 44.36mmol) and Nhydroxyphthalimide (7.24g, 44.36mmol). It was then dried 10 over P₂O₅ under high vacuum for two days at 40°C. The reaction mixture was flushed with argon and dry THF (369.8mL, Aldrich, sure seal bottle) was added to get a clear solution. Diethyl-azodicarboxylate (6.98mL, 44.36mmol) was added dropwise to the reaction mixture. The 15 rate of addition is maintained such that resulting deep red coloration is just discharged before adding the next drop. After the addition was complete, the reaction was stirred for 4 hrs. By that time TLC showed the completion of the reaction (ethylacetate:hexane, 60:40). The solvent was 20 evaporated in vacuum. Residue obtained was placed on a flash column and eluted with ethyl acetate:hexane (60:40), to get 2'-O-([2-phthalimidoxy)ethyl]-5'-tbutyldiphenylsilyl-5-methyluridine as white foam (21.819 g, 86%).

25

5

5'-O-tert-butyldiphenylsily1-2'-O-[(2-formadoximinooxy)ethyl]-5-methyluridine

2'-O-([2-phthalimidoxy)ethyl]-5'-t-butyldiphenylsilyl-5-methyluridine (3.1g, 4.5mmol) was dissolved in dry CH₂Cl₂
30 (4.5mL) and methylhydrazine (300mL, 4.64mmol) was added dropwise at -10°C to 0°C. After 1 h the mixture was filtered, the filtrate was washed with ice cold CH₂Cl₂ and the combined organic phase was washed with water, brine and dried over anhydrous Na₂SO₄. The solution was concentrated

10

to get 2'-0-(aminooxyethyl) thymidine, which was then
dissolved in MeOH (67.5mL). To this formaldehyde (20%
aqueous solution, w/w, 1.1 eq.) was added and the resulting
mixture was strirred for 1 h. Solvent was removed under
vacuum; residue chromatographed to get 5'-0-tertbutyldiphenylsilyl-2'-0-[(2-formadoximinooxy) ethyl]-5methyluridine as white foam (1.95 g, 78%).

5'-O-tert-Butyldiphenylsily1-2'-O-[N,N-dimethylaminooxyethyl]-5-methyluridine

5'-0-tert-butyldiphenylsilyl-2'-0-[(2formadoximinooxy)ethyl]-5-methyluridine (1.77q, 3.12mmol) was dissolved in a solution of 1M pyridinium ptoluenesulfonate (PPTS) in dry MeOH (30.6mL). Sodium 15 cyanoborohydride (0.39g, 6.13mmol) was added to this solution at 10°C under inert atmosphere. The reaction mixture was stirred for 10 minutes at 10°C. After that the reaction vessel was removed from the ice bath and stirred at room temperature for 2 h, the reaction monitored by TLC (5% MeOH in CH₂Cl₂). Aqueous NaHCO₃ solution (5%, 10mL) was 20 added and extracted with ethyl acetate (2x20mL). Ethyl acetate phase was dried over anhydrous Na₂SO₄, evaporated to dryness. Residue was dissolved in a solution of 1M PPTS in MeOH (30.6mL). Formaldehyde (20% w/w, 30mL, 3.37mmol) was 25 added and the reaction mixture was stirred at room temperature for 10 minutes. Reaction mixture cooled to 10°C in an ice bath, sodium cyanoborohydride (0.39g, 6.13mmol) was added and reaction mixture stirred at 10°C for 10 minutes. After 10 minutes, the reaction mixture was 30 removed from the ice bath and stirred at room temperature for 2 hrs. To the reaction mixture 5% NaHCO3 (25mL) solution was added and extracted with ethyl acetate (2x25mL). Ethyl acetate layer was dried over anhydrous Na_2SO_4 and evaporated to dryness . The residue obtained was

-62-

purified by flash column chromatography and eluted with 5% MeOH in CH₂Cl₂ to get 51-0-tert-butyldiphenylsilyl-21-0- [N,N-dimethylaminooxyethyl]-5-methyluridine as a white foam (14.6q, 80%).

5

2'-O-(dimethylaminooxyethyl)-5-methyluridine

Triethylamine trihydrofluoride (3.91mL, 24.0mmol) was dissolved in dry THF and triethylamine (1.67mL, 12mmol, dry, kept over KOH). This mixture of triethylamine-2HF was then added to 5'-O-tert-butyldiphenylsilyl-2'-O-[N,N-dimethylaminooxyethyl]-5-methyluridine (1.40g, 2.4mmol) and stirred at room temperature for 24 hrs. Reaction was monitored by TLC (5% MeOH in CH₂Cl₂). Solvent was removed under vacuum and the residue placed on a flash column and eluted with 10% MeOH in CH₂Cl₂ to get 2'-O-(dimethylaminooxyethyl)-5-methyluridine (766mg, 92.5%).

5'-O-DMT-2'-O-(dimethylaminooxyethyl)-5-methyluridine

2'-O-(dimethylaminooxyethyl)-5-methyluridine (750mg, 2.17mmol) was dried over P_2O_5 under high vacuum overnight at 40°C . It was then co-evaporated with anhydrous pyridine (20mL). The residue obtained was dissolved in pyridine (11mL) under argon atmosphere. 4-dimethylaminopyridine (26.5mg, 2.60mmol), 4,4'-dimethoxytrityl chloride (880mg,

2.60mmol) was added to the mixture and the reaction mixture was stirred at room temperature until all of the starting material disappeared. Pyridine was removed under vacuum and the residue chromatographed and eluted with 10% MeOH in CH₂Cl₂ (containing a few drops of pyridine) to get 5'-O-DMT-

30 2'-O-(dimethylamino-oxyethyl)-5-methyluridine (1.13g, 80%).

5'-O-DMT-2'-O-(2-N,N-dimethylaminooxyethyl)-5methyluridine-3'-[(2-cyanoethyl)-N,Ndiisopropylphosphoramidite]

5'-O-DMT-2'-O-(dimethylaminooxyethyl)-5-methyluridine 5 (1.08q, 1.67mmol) was co-evaporated with toluene (20mL). To the residue N, N-diisopropylamine tetrazonide (0.29g, 1.67mmol) was added and dried over P2O5 under high vacuum overnight at 40°C. Then the reaction mixture was dissolved in anhydrous acetonitrile (8.4mL) and 2-cyanoethyl-10 N,N,N¹,N¹-tetraisopropylphosphoramidite (2.12mL, 6.08mmol) was added. The reaction mixture was stirred at ambient temperature for 4 hrs under inert atmosphere. The progress of the reaction was monitored by TLC (hexane:ethyl acetate 1:1). The solvent was evaporated, then the residue was 15 dissolved in ethyl acetate (70mL) and washed with 5% aqueous NaHCO3 (40mL). Ethyl acetate layer was dried over anhydrous Na₂SO₄ and concentrated. Residue obtained was chromatographed (ethyl acetate as eluent) to get 5'-O-DMT-2'-O-(2-N, N-dimethylaminooxyethyl)-5-methyluridine-3'-[(2-20 cyanoethyl)-N,N-diisopropylphosphoramidite] as a foam (1.04g, 74.9%).

2'-(Aminooxyethoxy) nucleoside amidites

- 2'-(Aminooxyethoxy) nucleoside amidites [also known in the art as 2'-0-(aminooxyethyl) nucleoside amidites] are prepared as described in the following paragraphs.

 Adenosine, cytidine and thymidine nucleoside amidites are prepared similarly.
- N2-isobutyryl-6-O-diphenylcarbamoyl-2'-O-(2-ethylacetyl)-5'-O-(4,4'-dimethoxytrityl)guanosine-3'[(2-cyanoethyl)-N,N-diisopropylphosphoramidite]
 The 2'-O-aminooxyethyl guanosine analog may be

obtained by selective 2'-O-alkylation of diaminopurine

- riboside. Multigram quantities of diaminopurine riboside may be purchased from Schering AG (Berlin) to provide 2'-0-(2-ethylacetyl) diaminopurine riboside along with a minor amount of the 3'-0-isomer. 2'-0-(2-ethylacetyl)
- diaminopurine riboside may be resolved and converted to 2'-O-(2-ethylacetyl)guanosine by treatment with adenosine deaminase. (McGee, D. P. C., Cook, P. D., Guinosso, C. J., WO 94/02501 A1 940203.) Standard protection procedures should afford 2'-O-(2-ethylacetyl)-5'-O-(4,4'-
- dimethoxytrityl) guanosine and 2-N-isobutyryl-6-0-diphenylcarbamoyl-2'-O-(2-ethylacetyl)-5'-O-(4,4'-dimethoxytrityl) guanosine which may be reduced to provide 2-N-isobutyryl-6-O-diphenylcarbamoyl-2'-O-(2-ethylacetyl)-5'-O-(4,4'-dimethoxytrityl) guanosine. As before the
- hydroxyl group may be displaced by N-hydroxyphthalimide via a Mitsunobu reaction, and the protected nucleoside may phosphitylated as usual to yield 2-N-isobutyryl-6-0-diphenylcarbamoyl-2'-O-(2-ethylacetyl)-5'-O-(4,4'-dimethoxytrityl)guanosine-3'-[(2-cyanoethyl)-N,N-diisopropylphosphoramidite].

2'-dimethylaminoethoxyethoxy (2'-DMAEOE) nucleoside amidites

2'-dimethylaminoethoxyethoxy nucleoside amidites (also known in the art as 2'-O-dimethylaminoethoxyethyl, i.e., 2'-O-CH₂-O-CH₂-N(CH₂)₂, or 2'-DMAEOE nucleoside amidites) are prepared as follows. Other nucleoside amidites are prepared similarly.

30 2'-0-[2(2-N,N-dimethylaminoethoxy)ethyl]-5-methyl uridine

2[2-(Dimethylamino)ethoxy]ethanol (Aldrich, 6.66 g, 50 mmol) is slowly added to a solution of borane in tetrahydrofuran (1 M, 10 mL, 10 mmol) with stirring in a 100 mL bomb. Hydrogen gas evolves as the solid dissolves. O^2 -,2'-anhydro-5-methyluridine (1.2 g, 5 mmol), and sodium

WO 02/22635

bicarbonate (2.5 mg) are added and the bomb is sealed,

placed-in an oil-bath and heated to 155°C for 26 hours. The

bomb is cooled to room temperature and opened. The crude
solution is concentrated and the residue partitioned

between water (200 mL) and hexanes (200 mL). The excess
phenol is extracted into the hexane layer. The aqueous
layer is extracted with ethyl acetate (3x200 mL) and the
combined organic layers are washed once with water, dried
over anhydrous sodium sulfate and concentrated. The

residue is columned on silica gel using methanol/methylene

-65-

residue is columned on silica gel using methanol/methylene chloride 1:20 (which has 2% triethylamine) as the eluent. As the column fractions are concentrated a colorless solid forms which is collected to give the title compound as a white solid.

15

3.0

5'-O-dimethoxytrity1-2'-O-[2(2-N,N-dimethylaminoethoxy) ethyl)]-5-methyl uridine

To 0.5 g (1.3 mmol) of 2'-O-[2(2-N,N-dimethylamino-ethoxy)ethyl)]-5-methyl uridine in anhydrous pyridine (8 20 mL), triethylamine (0.36 mL) and dimethoxytrityl chloride (DMT-Cl, 0.87 g, 2 eq.) are added and stirred for 1 hour. The reaction mixture is poured into water (200 mL) and extracted with CH₂Cl₂ (2x200 mL). The combined CH₂Cl₂ layers are washed with saturated NaHCO₃ solution, followed by saturated NaCl solution and dried over anhydrous sodium sulfate. Evaporation of the solvent followed by silica gel chromatography using MeOH:CH₂Cl₂:Et₃N (20:1, v/v, with 1% triethylamine) gives the title compound.

5'-O-Dimethoxytrityl-2'-O-[2(2-N,N-dimethylaminoethoxy)ethyl)]-5-methyl uridine-3'-O-(cyanoethyl-N,N-diisopropyl)phosphoramidite

Diisopropylaminotetrazolide (0.6 g) and 2-cyanoethoxy-N,N-diisopropyl phosphoramidite (1.1 mL, 2 eq.) are added to a solution of 5'-O-dimethoxytrityl-2'-O-[2(2-N,N-dimethylaminoethoxy)ethyl)]-5-methyluridine (2.17-g, 3 mmol) dissolved in CH₂Cl₂ (20 mL) under an atmosphere of argon. The reaction mixture is stirred overnight and the solvent evaporated. The resulting residue is purified by silica gel flash column chromatography with ethyl acetate as the eluent to give the title compound.

Example 2

reference.

10 Oligonucleotide synthesis

Unsubstituted and substituted phosphodiester (P=O) oligonucleotides are synthesized on an automated DNA synthesizer (Applied Biosystems model 380B) using standard phosphoramidite chemistry with oxidation by iodine.

Phosphorothioates (P=S) are synthesized as for the phosphodiester oligonucleotides except the standard oxidation bottle was replaced by 0.2 M solution of 3H-1,2-benzodithiole-3-one 1,1-dioxide in acetonitrile for the stepwise thiation of the phosphite linkages. The thiation wait step was increased to 68 sec and was followed by the capping step. After cleavage from the CPG column and deblocking in concentrated ammonium hydroxide at 55°C (18 h), the oligonucleotides were purified by precipitating twice with 2.5 volumes of ethanol from a 0.5 M NaCl solution. Phosphinate oligonucleotides are prepared as described in U.S. Patent 5,508,270, herein incorporated by

Alkyl phosphonate oligonucleotides are prepared as described in U.S. Patent 4,469,863, herein incorporated by reference.

3'-Deoxy-3'-methylene phosphonate oligonucleotides are prepared as described in U.S. Patents 5,610,289 or 5,625,050, herein incorporated by reference.

Phosphoramidite oligonucleotides are prepared as described in U.S. Patent, 5,256,775 or U.S. Patent

-67-

5,366,878, herein incorporated by reference.

Alkylphosphonothioate oligonucleotides are prepared as described in published PCT applications PCT/US94/00902 and PCT/US93/06976 (published as WO 94/17093 and WO 94/02499, 5 respectively), herein incorporated by reference.

3'-Deoxy-3'-amino phosphoramidate oligonucleotides are prepared as described in U.S. Patent 5,476,925, herein incorporated by reference.

Phosphotriester oligonucleotides are prepared as described in U.S. Patent 5,023,243, herein incorporated by reference.

Borano phosphate oligonucleotides are prepared as described in U.S. Patents 5,130,302 and 5,177,198, both herein incorporated by reference.

15

30

35

Example 3

Oligonucleoside Synthesis

Methylenemethylimino linked oligonucleosides, also identified as MMI linked oligonucleosides, methylenedi20 methylhydrazo linked oligonucleosides, also identified as MDH linked oligonucleosides, and methylenecarbonylamino linked oligonucleosides, also identified as amide-3 linked oligonucleosides, and methyleneaminocarbonyl linked oligonucleosides, and methyleneaminocarbonyl linked oligonucleosides, also identified as amide-4 linked oligonucleosides, as well as mixed backbone compounds having, for instance, alternating MMI and P=O or P=S linkages are prepared as described in U.S. Patents 5,378,825, 5,386,023, 5,489,677, 5,602,240 and 5,610,289, all of which are herein incorporated by reference.

Formacetal and thioformacetal linked oligonucleosides are prepared as described in U.S. Patents 5,264,562 and 5,264,564, herein incorporated by reference.

Ethylene oxide linked oligonucleosides are prepared as described in U.S. Patent 5,223,618, herein incorporated by reference.

Example 4

PNA Synthesis

Peptide nucleic acids (PNAs) are prepared in

5 accordance with any of the various procedures referred to
in Peptide Nucleic Acids (PNA): Synthesis, Properties and
Potential Applications, Bioorganic & Medicinal Chemistry,
1996, 4, 5-23. They may also be prepared in accordance
with U.S. Patents 5,539,082, 5,700,922, and 5,719,262,
10 herein incorporated by reference.

Example 5

25

Synthesis of Chimeric Oligonucleotides

Chimeric oligonucleotides, oligonucleosides or mixed oligonucleotides/oligonucleosides of the invention can be of several different types. These include a first type wherein the "gap" segment of linked nucleosides is positioned between 5' and 3' "wing" segments of linked nucleosides and a second "open end" type wherein the "gap" segment is located at either the 3' or the 5' terminus of the oligomeric compound. Oligonucleotides of the first type are also known in the art as "gapmers" or gapped oligonucleotides. Oligonucleotides of the second type are also known in the art as "hemimers" or "wingmers".

[2'-O-Me] -- [2'-deoxy] -- [2'-O-Me] Chimeric Phosphorothicate Oligonucleotides

Chimeric oligonucleotides having 2'-O-alkyl phosphorothicate and 2'-deoxy phosphorothicate oligonucleotide segments are synthesized using an Applied

Biosystems automated DNA synthesizer Model 380B, as above. Oligonucleotides are synthesized using the automated synthesizer and 2'-deoxy-5'-dimethoxytrityl-3'-O-phosphoramidite for the DNA portion and 5'-dimethoxytrityl-2'-O-methyl-3'-O-phosphoramidite for 5' and 3' wings. The

-69-

standard synthesis cycle is modified by increasing the wait step after the delivery of tetrazole and base to 600 s repeated four times for RNA and twice for 2'-O-methyl. fully protected oligonucleotide is cleaved from the support 5 and the phosphate group is deprotected in 3:1 ammonia/ethanol at room temperature overnight then lyophilized to dryness. Treatment in methanolic ammonia for 24 hrs at room temperature is then done to deprotect all bases and sample was again lyophilized to dryness. 10 pellet is resuspended in 1M TBAF in THF for 24 hrs at room temperature to deprotect the 2' positions. The reaction is then quenched with 1M TEAA and the sample is then reduced to 1/2 volume by rotovac before being desalted on a G25 size exclusion column. The oligo recovered is then analyzed spectrophotometrically for yield and for purity by capillary electrophoresis and by mass spectrometry.

[2'-O-(2-Methoxyethyl)]--[2'-deoxy]--[2'-O-(Methoxyethyl)] Chimeric Phosphorothicate Oligonucleotides

20

25

[2'-O-(2-methoxyethyl)]--[2'-deoxy]--[-2'-O-(methoxyethyl)] chimeric phosphorothioate oligonucleotides were prepared as per the procedure above for the 2'-O-methyl chimeric oligonucleotide, with the substitution of 2'-O-(methoxyethyl) amidites for the 2'-O-methyl amidites.

[2'-0-(2-Methoxyethyl) Phosphodiester] -- [2'-deoxy Phosphorothioate] -- [2'-0-(2-Methoxyethyl) Phosphodiester] Chimeric Oligonucleotides

2'-O-(2-methoxyethyl phosphodiester]--[2'-deoxy phosphorothioate]--[2'-O-(methoxyethyl) phosphodiester]

chimeric oligonucleotides are prepared as per the above procedure for the 2'-O-methyl chimeric oligonucleotide with the substitution of 2'-O-(methoxyethyl) amidites for the

2'-O-methyl amidites, oxidization with iodine to generate

-70-

the phosphodiester internucleotide linkages within the

wing portions of the chimeric structures and sulfurization
utilizing 3,H-1,2 benzodithiole-3-one 1,1 dioxide (Beaucage
Reagent) to generate the phosphorothioate internucleotide

5 linkages for the center gap.

Other chimeric oligonucleotides, chimeric oligonucleosides and mixed chimeric oligonucleotides/oligonucleosides are synthesized according to United States patent 5,623,065, herein incorporated by reference.

10

Example 6

Oligonucleotide Isolation

After cleavage from the controlled pore glass column (Applied Biosystems) and deblocking in concentrated 15 ammonium hydroxide at 55°C for 18 hours, the oligonucleotides or oligonucleosides are purified by precipitation twice out of 0.5 M NaCl with 2.5 volumes ethanol. Synthesized oligonucleotides were analyzed by polyacrylamide gel electrophoresis on denaturing gels and 20 judged to be at least 85% full length material. relative amounts of phosphorothicate and phosphodiester linkages obtained in synthesis were periodically checked by ³¹P nuclear magnetic resonance spectroscopy, and for some studies oligonucleotides were purified by HPLC, as 25 described by Chiang et al., J. Biol. Chem. 1991, 266, 18162-18171. Results obtained with HPLC-purified material were similar to those obtained with non-HPLC purified material.

30 Example 7

Oligonucleotide Synthesis - 96 Well Plate Format

Oligonucleotides were synthesized via solid phase P(III) phosphoramidite chemistry on an automated synthesizer capable of assembling 96 sequences

-71-

simultaneously in a standard 96 well format.

Phosphodiester internucleotide linkages were afforded by oxidation with aqueous iodine. Phosphorothioate internucleotide linkages were generated by sulfurization

tilizing 3,H-1,2 benzodithiole-3-one 1,1 dioxide (Beaucage Reagent) in anhydrous acetonitrile. Standard base-protected beta-cyanoethyldiisopropyl phosphoramidites were purchased from commercial vendors (e.g. PE-Applied Biosystems, Foster City, CA, or Pharmacia, Piscataway, NJ).

Non-standard nucleosides are synthesized as per known literature or patented methods. They are utilized as base protected beta-cyanoethyldiisopropyl phosphoramidites.

Oligonucleotides were cleaved from support and deprotected with concentrated NH₄OH at elevated temperature (55-60°C) for 12-16 hours and the released product then dried in vacuo. The dried product was then re-suspended in sterile water to afford a master plate from which all analytical and test plate samples are then diluted utilizing robotic pipettors.

20

Example 8

Oligonucleotide Analysis - 96 Well Plate Format

The concentration of oligonucleotide in each well was assessed by dilution of samples and UV absorption

25 spectroscopy. The full-length integrity of the individual products was evaluated by capillary electrophoresis (CE) in either the 96 well format (Beckman P/ACETM MDQ) or, for individually prepared samples, on a commercial CE apparatus (e.g., Beckman P/ACETM 5000, ABI 270). Base and backbone

30 composition was confirmed by mass analysis of the compounds utilizing electrospray-mass spectroscopy. All assay test plates were diluted from the master plate using single and multi-channel robotic pipettors. Plates were judged to be acceptable if at least 85% of the compounds on the plate

-72-

were at least 85% full length.

Example 9

Cell culture and oligonucleotide treatment

The effect of antisense compounds on target nucleic acid expression can be tested in any of a variety of cell types provided that the target nucleic acid is present at measurable levels. This can be routinely determined using, for example, PCR or Northern blot analysis. The following 4 cell types are provided for illustrative purposes, but other cell types can be routinely used, provided that the target is expressed in the cell type chosen. This can be readily determined by methods routine in the art, for example Northern blot analysis, Ribonuclease protection assays, or RT-PCR.

T-24 cells:

The human transitional cell bladder carcinoma cell line T-24 was obtained from the American Type Culture

Collection (ATCC) (Manassas, VA). T-24 cells were routinely cultured in complete McCoy's 5A basal media (Gibco/Life Technologies, Gaithersburg, MD) supplemented with 10% fetal calf serum (Gibco/Life Technologies, Gaithersburg, MD), penicillin 100 units per mL, and

streptomycin 100 micrograms per mL (Gibco/Life Technologies, Gaithersburg, MD). Cells were routinely passaged by trypsinization and dilution when they reached 90% confluence. Cells were seeded into 96-well plates (Falcon-Primaria #3872) at a density of 7000 cells/well for use in RT-PCR analysis.

For Northern blotting or other analysis, cells may be seeded onto 100 mm or other standard tissue culture plates and treated similarly, using appropriate volumes of medium and oligonucleotide.

-73-

A549 cells:

The human lung carcinoma cell line A549 was obtained from the American Type Culture Collection (ATCC) (Manassas, VA). A549 cells were routinely cultured in DMEM basal media (Gibco/Life Technologies, Gaithersburg, MD) supplemented with 10% fetal calf serum (Gibco/Life Technologies, Gaithersburg, MD), penicillin 100 units per mL, and streptomycin 100 micrograms per mL (Gibco/Life Technologies, Gaithersburg, MD). Cells were routinely passaged by trypsinization and dilution when they reached 90% confluence.

NHDF cells:

Human neonatal dermal fibroblast (NHDF) were obtained from the Clonetics Corporation (Walkersville MD). NHDFs were routinely maintained in Fibroblast Growth Medium (Clonetics Corporation, Walkersville MD) supplemented as recommended by the supplier. Cells were maintained for up to 10 passages as recommended by the supplier.

20

HEK cells:

Human embryonic keratinocytes (HEK) were obtained from the Clonetics Corporation (Walkersville MD). HEKs were routinely maintained in Keratinocyte Growth Medium

25 (Clonetics Corporation, Walkersville MD) formulated as recommended by the supplier. Cells were routinely maintained for up to 10 passages as recommended by the supplier.

30 Treatment with antisense compounds:

When cells reached 80% confluency, they were treated with oligonucleotide. For cells grown in 96-well plates, wells were washed once with 200 μ L OPTI-MEMTM-1 reduced-serum medium (Gibco BRL) and then treated with 130 μ L of

OPTI-MEMTM-1 containing 3.75 μg/mL LIPOFECTINTM (Gibco BRL)

and the desired concentration of oligonucleotide. After-4---
7 hours of treatment, the medium was replaced with fresh medium. Cells were harvested 16-24 hours after

5 oligonucleotide treatment.

The concentration of oligonucleotide used varies from cell line to cell line. To determine the optimal oligonucleotide concentration for a particular cell line, the cells are treated with a positive control

- oligonucleotide at a range of concentrations. For human cells the positive control oligonucleotide is ISIS 13920,

 TCCGTCATCGCTCCTCAGGG, SEQ ID NO: 1, a 2'-O-methoxyethyl gapmer (2'-O-methoxyethyls shown in bold) with a phosphorothioate backbone which is targeted to human H-ras.
- 15 For mouse or rat cells the positive control oligonucleotide is ISIS 15770, ATGCATTCTGCCCCCAAGGA, SEQ ID NO: 2, a 2'-O-methoxyethyl gapmer (2'-O-methoxyethyls shown in bold) with a phosphorothicate backbone which is targeted to both mouse and rat c-raf. The concentration of positive
- control oligonucleotide that results in 80% inhibition of c-Ha-ras (for ISIS 13920) or c-raf (for ISIS 15770) mRNA is then utilized as the screening concentration for new oligonucleotides in subsequent experiments for that cell line. If 80% inhibition is not achieved, the lowest
- concentration of positive control oligonucleotide that results in 60% inhibition of H-ras or c-raf mRNA is then utilized as the oligonucleotide screening concentration in subsequent experiments for that cell line. If 60% inhibition is not achieved, that particular cell line is
- 30 deemed as unsuitable for oligonucleotide transfection experiments.

-75-

Example 10

Analysis of oligonucleotide inhibition of clusterin expression

Antisense modulation of clusterin expression can be 5 assayed in a variety of ways known in the art. For example, clusterin mRNA levels can be quantitated by, e.g., Northern blot analysis, competitive polymerase chain reaction (PCR), or real-time PCR (RT-PCR). Real-time quantitative PCR is presently preferred. RNA analysis can 10 be performed on total cellular RNA or poly(A) + mRNA. Methods of RNA isolation are taught in, for example, Ausubel, F.M. et al., Current Protocols in Molecular Biology, Volume 1, pp. 4.1.1-4.2.9 and 4.5.1-4.5.3, John Wiley & Sons, Inc., 1993. Northern blot analysis is 15 routine in the art and is taught in, for example, Ausubel, F.M. et al., Current Protocols in Molecular Biology, Volume 1, pp. 4.2.1-4.2.9, John Wiley & Sons, Inc., 1996. Realtime quantitative (PCR) can be conveniently accomplished using the commercially available ABI PRISMTM 7700 Sequence 20 Detection System, available from PE-Applied Biosystems, Foster City, CA and used according to manufacturer's instructions. Prior to quantitative PCR analysis, primerprobe sets specific to the target gene being measured are evaluated for their ability to be "multiplexed" with a 25 GAPDH amplification reaction. In multiplexing, both the target gene and the internal standard gene GAPDH are amplified concurrently in a single sample. In this analysis, mRNA isolated from untreated cells is serially diluted. Each dilution is amplified in the presence of 30 primer-probe sets specific for GAPDH only, target gene only ("single-plexing"), or both (multiplexing). Following PCR amplification, standard curves of GAPDH and target mRNA signal as a function of dilution are generated from both the single-plexed and multiplexed samples. If both the

slope and correlation coefficient of the GAPDH and target
signals generated from the multiplexed samples fall within
10% of their corresponding values generated from the
single-plexed samples, the primer-probe set specific for
that target is deemed as multiplexable. Other methods of
PCR are also known in the art.

Protein levels of clusterin can be quantitated in a variety of ways well known in the art, such as immunoprecipitation, Western blot analysis

- 10 (immunoblotting), ELISA or fluorescence-activated cell sorting (FACS). Antibodies directed to clusterin can be identified and obtained from a variety of sources, such as the MSRS catalog of antibodies (Aerie Corporation, Birmingham, MI), or can be prepared via conventional
- antibody generation methods. Methods for preparation of polyclonal antisera are taught in, for example, Ausubel, F.M. et al., Current Protocols in Molecular Biology, Volume 2, pp. 11.12.1-11.12.9, John Wiley & Sons, Inc., 1997. Preparation of monoclonal antibodies is taught in, for
- 20 example, Ausubel, F.M. et al., Current Protocols in
 Molecular Biology, Volume 2, pp. 11.4.1-11.11.5, John Wiley
 & Sons, Inc., 1997.

Immunoprecipitation methods are standard in the art and can be found at, for example, Ausubel, F.M. et al.,

- 25 Current Protocols in Molecular Biology, Volume 2, pp. 10.16.1-10.16.11, John Wiley & Sons, Inc., 1998. Western blot (immunoblot) analysis is standard in the art and can be found at, for example, Ausubel, F.M. et al., Current Protocols in Molecular Biology, Volume 2, pp. 10.8.1-
- 30 10.8.21, John Wiley & Sons, Inc., 1997. Enzyme-linked immunosorbent assays (ELISA) are standard in the art and can be found at, for example, Ausubel, F.M. et al., Current Protocols in Molecular Biology, Volume 2, pp. 11.2.1-11.2.22, John Wiley & Sons, Inc., 1991.

Example 11

Poly(A) + mRNA isolation

Poly(A) + mRNA was isolated according to Miura et al., Clin. Chem., 1996, 42, 1758-1764. Other methods for poly(A) + mRNA isolation are taught in, for example, Ausubel, F.M. et al., Current Protocols in Molecular Biology, Volume 1, pp. 4.5.1-4.5.3, John Wiley & Sons, Inc., 1993. Briefly, for cells grown on 96-well plates, growth medium was removed from the cells and each well was washed with 200 μL cold PBS. 60 μL lysis buffer (10 mM Tris-HCl, pH 7.6, 1 mM EDTA, 0.5 M NaCl, 0.5% NP-40, 20 mM vanadyl-ribonucleoside complex) was added to each well, the plate was gently agitated and then incubated at room 15 temperature for five minutes. 55 μL of lysate was transferred to Oligo d(T) coated 96-well plates (AGCT Inc., Irvine CA). Plates were incubated for 60 minutes at room temperature, washed 3 times with 200 µL of wash buffer (10 mM Tris-HCl pH 7.6, 1 mM EDTA, 0.3 M NaCl). After the 20 final wash, the plate was blotted on paper towels to remove excess wash buffer and then air-dried for 5 minutes. $60 \mu L$ of elution buffer (5 mM Tris-HCl pH 7.6), preheated to 70°C was added to each well, the plate was incubated on a 90°C hot plate for 5 minutes, and the eluate was then

Cells grown on 100 mm or other standard plates may be treated similarly, using appropriate volumes of all solutions.

30 Example 12

Total RNA Isolation

25 transferred to a fresh 96-well plate.

Total mRNA was isolated using an RNEASY 96™ kit and buffers purchased from Qiagen Inc. (Valencia CA) following

-78-

the manufacturer's recommended procedures. Briefly, for cells grown on 96-well plates, growth medium was removed from the cells and each well was washed with 200 µL cold PBS. 100 μ L Buffer RLT was added to each well and the plate 5 vigorously agitated for 20 seconds. 100 μL of 70% ethanol was then added to each well and the contents mixed by pipetting three times up and down. The samples were then transferred to the RNEASY 96^{TM} well plate attached to a QIAVACTM manifold fitted with a waste collection tray and 10 attached to a vacuum source. Vacuum was applied for 15 seconds. 1 mL of Buffer RW1 was added to each well of the RNEASY 96™ plate and the vacuum again applied for 15 seconds. 1 mL of Buffer RPE was then added to each well of the RNEASY 96TM plate and the vacuum applied for a period of 15 15 seconds. The Buffer RPE wash was then repeated and the vacuum was applied for an additional 10 minutes. The plate was then removed from the QIAVAC TM manifold and blotted dry on paper towels. The plate was then re-attached to the QIAVACTM manifold fitted with a collection tube rack 20 containing 1.2 mL collection tubes. RNA was then eluted by pipetting 60 µL water into each well, incubating 1 minute, and then applying the vacuum for 30 seconds. The elution step was repeated with an additional 60 µL water.

The repetitive pipetting and elution steps may be
25 automated using a QIAGEN Bio-Robot 9604 (Qiagen, Inc.,
Valencia CA). Essentially, after lysing of the cells on
the culture plate, the plate is transferred to the robot
deck where the pipetting, DNase treatment and elution steps
are carried out.

Example 13

Real-time Quantitative PCR Analysis of clusterin mRNA Levels

Quantitation of clusterin mRNA levels was determined 5 by real-time quantitative PCR using the ABI PRISM™ 7700 Sequence Detection System (PE-Applied Biosystems, Foster City, CA) according to manufacturer's instructions. is a closed-tube, non-gel-based, fluorescence detection system which allows high-throughput quantitation of 10 polymerase chain reaction (PCR) products in real-time. As opposed to standard PCR, in which amplification products are quantitated after the PCR is completed, products in real-time quantitative PCR are quantitated as they accumulate. This is accomplished by including in the PCR 15 reaction an oligonucleotide probe that anneals specifically between the forward and reverse PCR primers, and contains two fluorescent dyes. A reporter dye (e.g., JOE, FAM, or VIC, obtained from either Operon Technologies Inc., Alameda, CA or PE-Applied Biosystems, Foster City, CA) is 20 attached to the 5' end of the probe and a quencher dye (e.g., TAMRA, obtained from either Operon Technologies Inc., Alameda, CA or PE-Applied Biosystems, Foster City, CA) is attached to the 3' end of the probe. When the probe and dyes are intact, reporter dye emission is quenched by 25 the proximity of the 3' quencher dye. During amplification, annealing of the probe to the target sequence creates a substrate that can be cleaved by the 5'exonuclease activity of Taq polymerase. During the extension phase of the PCR amplification cycle, cleavage of 30 the probe by Taq polymerase releases the reporter dye from the remainder of the probe (and hence from the quencher moiety) and a sequence-specific fluorescent signal is generated. With each cycle, additional reporter dye molecules are cleaved from their respective probes, and the 35 fluorescence intensity is monitored at regular intervals by

laser optics built into the ABI PRISMTM 7700 Sequence

Detection System. In each assay, a series of parallel

reactions containing serial dilutions of mRNA from

untreated control samples generates a standard curve that

is used to quantitate the percent inhibition after

antisense oligonucleotide treatment of test samples.

PCR reagents were obtained from PE-Applied Biosystems, Foster City, CA. RT-PCR reactions were carried out by adding 25 μL PCR cocktail (1x TAQMANTM buffer A, 5.5 mM 10 MgCl₂, 300 μM each of dATP, dCTP and dGTP, 600 μM of dUTP, 100 nM each of forward primer, reverse primer, and probe, 20 Units RNAse inhibitor, 1.25 Units AMPLITAQ GOLDTM, and 12.5 Units MuLV reverse transcriptase) to 96 well plates containing 25 μL poly(A) mRNA solution. The RT reaction 15 was carried out by incubation for 30 minutes at 48°C. Following a 10 minute incubation at 95°C to activate the AMPLITAQ GOLDTM, 40 cycles of a two-step PCR protocol were carried out: 95°C for 15 seconds (denaturation) followed by 60°C for 1.5 minutes (annealing/extension).

- Probes and primers to human clusterin were designed to hybridize to a human clusterin sequence, using published sequence information (GenBank accession number M64722, incorporated herein as SEQ ID NO:3). For human clusterin the PCR primers were:
- forward primer: TCCGTACGAGCCCCTGAA (SEQ ID NO: 4)
 reverse primer: TGAGCCTCGTGTATCATCTCAAG (SEQ ID NO: 5) and
 the PCR probe was: FAM-TCCACGCCATGTTCCAGCCCT-TAMRA
 (SEQ ID NO: 6) where FAM (PE-Applied Biosystems, Foster
 City, CA) is the fluorescent reporter dye) and TAMRA (PE-
- 30 Applied Biosystems, Foster City, CA) is the quencher dye. For human GAPDH the PCR primers were:

forward primer: CAACGGATTTGGTCGTATTGG (SEQ ID NO: 7)
reverse primer: GGCAACAATATCCACTTTACCAGAGT (SEQ ID NO: 8)

-81-

and the PCR probe was: 5' JOE-CGCCTGGTCACCAGGGCTGCT- TAMRA 3! (SEQ_ID_NO: 9) where JOE_(PE-Applied_Biosystems, Foster_ City, CA) is the fluorescent reporter dye) and TAMRA (PE-Applied Biosystems, Foster City, CA) is the quencher dye.

5

25

Example 14

Northern blot analysis of clusterin mRNA levels

Eighteen hours after antisense treatment, cell monolayers were washed twice with cold PBS and lysed in 1 10 mL RNAZOLTM (TEL-TEST "B" Inc., Friendswood, TX). Total RNA was prepared following manufacturer's recommended protocols. Twenty micrograms of total RNA was fractionated by electrophoresis through 1.2% agarose gels containing 1.1% formaldehyde using a MOPS buffer system (AMRESCO, Inc. 15 Solon, OH). RNA was transferred from the gel to HYBONDTM-N+ nylon membranes (Amersham Pharmacia Biotech, Piscataway, NJ) by overnight capillary transfer using a Northern/Southern Transfer buffer system (TEL-TEST "B" Inc., Friendswood, TX). RNA transfer was confirmed by UV 20 visualization. Membranes were fixed by UV cross-linking using a STRATALINKERTM UV Crosslinker 2400 (Stratagene, Inc, La Jolla, CA) and then robed using QUICKHYBTM hybridization solution (Stratagene, La Jolla, CA) using

To detect human clusterin, a human clusterin specific probe was prepared by PCR using the forward primer TCCGTACGAGCCCCTGAA (SEQ ID NO: 4) and the reverse primer TGAGCCTCGTGTATCATCTCAAG (SEQ ID NO: 5). To normalize for variations in loading and transfer efficiency membranes 30 were stripped and probed for human glyceraldehyde-3phosphate dehydrogenase (GAPDH) RNA (Clontech, Palo Alto, CA).

manufacturer's recommendations for stringent conditions.

Hybridized membranes were visualized and quantitated using a PHOSPHORIMAGER™ and IMAGEQUANT™ Software V3.3

(Molecular Dynamics, Sunnyvale, CA). Data was normalized to GAPDH levels in untreated controls.

Example 15

5 Antisense inhibition of human clusterin expression by chimeric phosphorothicate oligonucleotides having 2'-MOE wings and a deoxy gap

In accordance with the present invention, a series of oligonucleotides were designed to target different regions 10 of the human clusterin RNA, using published sequences (GenBank accession number M64722, incorporated herein as SEQ ID NO: 3, GenBank accession number L00974, incorporated herein as SEQ ID NO: 10, GenBank accession number M63377, incorporated herein as SEQ ID NO: 11, GenBank accession 15 number M63376, incorporated herein as SEQ ID NO: 12, and GenBank accession number M25915, incorporated herein as SEQ ID NO: 13). The oligonucleotides are shown in Table 1. "Target site" indicates the first (5'-most) nucleotide number on the particular target sequence to which the 20 oligonucleotide binds. All compounds in Table 1 are chimeric oligonucleotides ("gapmers") 20 nucleotides in length, composed of a central "gap" region consisting of ten 2'-deoxynucleotides, which is flanked on both sides (5' and 3' directions) by five-nucleotide "wings". The wings 25 are composed of 2'-methoxyethyl (2'-MOE) nucleotides. internucleoside (backbone) linkages are phosphorothioate (P=S) throughout the oligonucleotide. All cytidine residues are 5-methylcytidines. The compounds were analyzed for their effect on human clusterin mRNA levels by 30 quantitative real-time PCR as described in other examples herein. Data are averages from two experiments. If present, "N.D." indicates "no data".

5

_____Table 1 _____

Inhibition of human clusterin mRNA levels by chimeric phosphorothicate oligonucleotides having 2'-MOE wings and a deoxy gap

ISIS #	REGION TARGET SEQ ID NO		TARGET SITE	SEQUENCE	%INHIB	SEQ ID NO
129045	5'UTR	3	18	gtctttgcacgcctcggtca	64	14
129046	5'UTR	3	26	attctggagtctttgcacgc	64	15
129047	Start	3	44	gtcttcatcatgcctccaat	68	16
	Codon					
129048	Coding	3	82	tctcccaggtcagcagcagc	67	17
129049	Coding	3	106	tetggteeeceaggaeetge	46	18
129050	Coding	3	127	ggagctcattgtctgagacc	59	19
129052	Coding	3	154	acttacttccctgattggac	77	20
129053	Coding	3	171	aatttccttattgacgtact	68	21
129054	Coding	3	206	gtctttatctgtttcacccc	85	. 22
129055	Coding	3	286	gggcatcctctttcttcttc	64	23
129056	Coding	3	291	atttagggcatcctcttct	31	24
129057	Coding	3	303	ttccctggtctcatttaggg	68	25
129058	Coding	3	312	tgtctctgattccctggtct	79	26
129059	Coding	3	329	gggagctccttcagctttgt	49	27
129060	Coding	3	364	cccagagggccatcatggtc	45	28
129061	Coding	3	369	ctcttcccagagggccatca	36	29
129062	Coding	3	385	tcaggcagggcttacactct	70	30
129063	Coding	3	412	gtgcgtagaacttcatgcag	60	31
129064	Coding	3	448	ggcggccaaccaggcctgag	42	32
129065	Coding	3	449	tggcggccaaccaggcctga	32	33
129066	Coding	3	460	actcctcaagctggcggcca	67	34
129067	Coding	3	487	agtagaaggggagctctgg	63	35
129068	Coding	3	497	ttcatccagaagtagaaggg	41	36
129069	Coding	3	522	cagcaggagtcgatgcggt	60	37
129070	Coding	3	538	gctgccggtcgttctccagc	51	38
129071	Coding	3	556	catccagcatgtgcgtctgc	69	39
129072	Coding	3	558	gacatccagcatgtgcgtct	55	40
129073	Coding	3	570	gtggtcctgcatgacatcca	62	41
129074	Coding	3	572	aagtggtcctgcatgacatc	41	42
129075	Coding	3	609	ctggaagagctcgtctatga	67	43
129076	Coding	3	613	tgtcctggaagagctcgtct	69	44
129077	Coding	3	618	gaacctgtcctggaagagct	68	45
129078	Coding	3	695	ggaaagaagaagtgaggcct	44	46
129079	Coding	3	726	gggcatcaagctgcggacga	65	47
129080	Coding	3	780	ctcaaggaaggctggaaca	81	48
129081	Coding	3	781	tctcaaggaagggctggaac	81	49
129082	Coding	3	788	tgtatcatctcaaggaaggg	38	50
129083	Coding	3	825	gctgtggaagtggatgtcca	48	51
129084	Coding	3	853	attctgttggcgggtgctgg	50	52
129085	Coding	3	858	tatgaattctgttggcgggt	36	53
129086	Coding	3	898	ggatctcccggcacacagtc	68	54
129087	Coding	3	899	cggatctcccggcacacagt	70	55

120000	G = 12 1		011		69	56
129088	Coding	3	911	gtggagttgtggcggatctc		
129089	Coding	3	933	gtccttcatccgcaggcagc	56	57
129090	Coding	3	972	acagtccacagacaagatct	49	58
129092	Coding	3	1014	gagetecegeegeagettag	22	59
129093	Coding	3	1027	ggagggattcgtcgagctcc	55	60
129094	Coding	3	1088	atcttccactggtaggactt	50	61
129095	Coding	3	1096	tgttgagcatcttccactgg	62	62
129096	Coding	3	1118	agctgctccagcaaggagga	46	63
129097	Coding	3	1126	gctcgttcagctgctccagc	43	64
129098	Coding	3	1153	ttgccagccgggacacccag	72	65
129099	Coding	3	1187	cgcagatagtactggtcttc	61	66
129100	Coding	3	1199	accgtggtgacccgcagata	73	67
129101	Coding	3	1221	cgagtcagaagtgtgggaag	24	68
129102	Coding	3	1280	gtgatgggatcagagtcaaa	38	69
129103	Coding	3	1305	ggagacttctacagggaccg	63	70
129104	Coding	3	1337	gccacggtctccataaattt	70	71
129106	3 'UTR	3	1403	gcaaaagcaacatccacatc	74	72
129107	3 'UTR	3	1550	tagagtgcaggatccagagc	71	73
129108	3 'UTR	3	1605	attagttgcatgcaggagca	71	74
129109	3 'UTR	3	1620	agacagttttattgaattag	11	75
129118	Intron	10	2819	cgagatagagccactgtacg	44	76
129119	Intron	10	4646	tgccaccacccccgggtgat	13	77
129091	Intron-	10	5849	gttgttggtggaacagtcca	40	78
	Exon					
	Junction					
129120	Intron-	10	7384	tgcttaccggtgctttttgc	46	79
	Exon					
	Junction					
129105	Intron-	10	7600	acatctcactcctcccggtg	70	80
	Exon					
	Junction				-	
129121	3'UTR	10	7855	gaccctccaagcgatcagct	20	81
129122	3'UTR	10	7863	aaaaagaggaccctccaagc	39	82
129115	Intron-	11	322	tgtgtccccttttcacctgg	54	83
	Exon	l				
	Junction					
129116	Intron	11	445	attaccaatggagcatggca	43	84
129117	Intron	11	810	caacatggccaaaccccatg	55	85
129112	Intron	12	1766	gcggcaggtctccaggtctc	43	86
129110	Intron	12	4813	ttcccttcggagagtagaga	44	87
129113	Intron	12	5848	tgcttgggaaatgcctgcaa	34	88
129114	Intron	12	6936	agctggatgccagaaaggcc	40	89
129111	5'UTR	13	39	tggaagtagtggaagccagg	11	90

As shown in Table 1, SEQ ID NOs 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 60, 61, 62, 63, 64, 65, 66, 67, 69, 70, 71, 72, 73, 74, 76, 78, 79, 80, 82, 83, 84, 85, 86, 87, 88 and 89 demonstrated at least 30%

-85-

inhibition of human clusterin expression in this assay and
are therefore preferred. The target sites to which these
preferred sequences are complementary are herein referred
to as "active sites" and are therefore preferred sites for
targeting by compounds of the present invention.

Example 16

Western blot analysis of clusterin protein levels

Western blot analysis (immunoblot analysis) is carried out using standard methods. Cells are harvested 16-20 h after oligonucleotide treatment, washed once with PBS, suspended in Laemmli buffer (100 ul/well), boiled for 5 minutes and loaded on a 16% SDS-PAGE gel. Gels are run for 1.5 hours at 150 V, and transferred to membrane for western blotting. Appropriate primary antibody directed to clusterin is used, with a radiolabelled or fluorescently labeled secondary antibody directed against the primary antibody species. Bands are visualized using a PHOSPHORIMAGERTM (Molecular Dynamics, Sunnyvale CA).

What is claimed is:

- 1. A compound 8 to 50 nucleobases in length which is targeted to the 3' UTR, an intron, an intron-exon junction, or nucleobases 106-1402 of the coding region of a nucleic acid molecule encoding clusterin, wherein said compound specifically hybridizes with and inhibits the expression of clusterin.
 - 2. The compound of claim 1 which is an antisense oligonucleotide.
 - 3. The compound of claim 2 wherein the antisense oligonucleotide has a sequence comprising SEQ ID NO: 18,
- 15 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33,
- 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48,
 - 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 60, 61, 62, 63, 64,
 - 65, 66, 67, 69, 70, 71, 72, 73, 74, 76, 78, 79, 80, 82, 83,
 - 84, 85, 86, 87, 88 or 89.
- 20 4. The compound of claim 2 wherein the antisense oligonucleotide comprises at least one modified internucleoside linkage.
 - 5. The compound of claim 4 wherein the modified internucleoside linkage is a phosphorothicate linkage.
- 25 6. The compound of claim 2 wherein the antisense oligonucleotide comprises at least one modified sugar moiety.
 - 7. The compound of claim 6 wherein the modified sugar moiety is a 2'-O-methoxyethyl sugar moiety.
- 30 8. The compound of claim 2 wherein the antisense oligonucleotide comprises at least one modified nucleobase.
 - 9. The compound of claim 8 wherein the modified nucleobase is a 5-methylcytosine.
- 10. The compound of claim 2 wherein the antisense 35 oligonucleotide is a chimeric oligonucleotide.

- 11. A compound 8 to 50 nucleobases in length which specifically hybridizes with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding clusterin.
 - 5 12. A composition comprising the compound of claim 1 and a pharmaceutically acceptable carrier or diluent.
 - 13. The composition of claim 12 further comprising a colloidal dispersion system.
 - 14. The composition of claim 12 wherein the compound 10 is an antisense oligonucleotide.
 - 15. A method of inhibiting the expression of clusterin in cells or tissues comprising contacting said cells or tissues with the compound of claim 1 so that expression of clusterin is inhibited.
 - 16. A method of treating an animal having a disease or condition associated with clusterin comprising administering to said animal a therapeutically or prophylactically effective amount of the compound of claim 1 so that expression of clusterin is inhibited.
 - 20 17. The method of claim 16 wherein the disease or condition is a hypercholesterolemia.
 - 18. The method of claim 16 wherein the disease or condition is a cardiovascular disorder.
 - 19. The method of claim 16 wherein the disease or 25 condition is a hyperproliferative disorder.
 - 20. The method of claim 16 wherein the disease or condition is a hyperlipidemic disorder.

SEQUENCE LISTING

```
<110> Isis Pharmaceuticals, Inc.
   Brett P. Monia
     Susan M. Freier
<120> ANTISENSE MODULATION OF CLUSTERIN EXPRESSION
<130> RTSP-0177
<150> 09/659,791
<151> 2000-09-11
<160> 90
<210> 1
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Antisense Oligonucleotide
<400> 1
tccgtcatcg ctcctcaggg
                                                                     20
<210> 2
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Antisense Oligonucleotide
<400> 2
atgcattctg cccccaagga
                                                                     20
<210> 3
<211> 1648
<212> DNA
<213> Homo sapiens
<220>
<221> CDS
<222> (53)...(1402)
<400> 3
cgcggacagg gtgccgctga ccgaggcgtg caaagactcc agaattggag gc atg atg
                                                          Met Met
aag act ctg ctg ctg ttt gtg ggg ctg ctg ctg acc tgg gag agt ggg
```

Lys	Thr	Leu 5	Leu	Leu	Phe	Val	Gly 10	Leu	Leu	Leu	Thr	Trp 15	Glu	Ser	Gly		
		_		_	_	_	_		_				_	gaa Glu			154
	aat	_		_	_	tac	_		_	_	att			gct Ala	_	;	202
				_		_				~				gaa Glu 65		:	250
_	_		_		_			-	_	_	_	_	_	aaa Lys		•	298
_	_						_					-	_	gag Glu		:	346
														tgt Cys		:	394
	_	_		_		_	-	_			-	_	-	tgc Cys		•	442
_				_	_		_	_					_	aac Asn 145	_	•	490
										_				tcc Ser		!	538
														cag Gln		!	586
														gac Asp		•	634
						_	_					_		ttc Phe	_	(682
_													_	atc Ile 225	_	•	730
cgc	agc	ttg	atg	ccc	ttc	tct	ccg	tac	gag 2	ccc	ctg	aac	ttc	cac	gcc	•	778

Arg	Ser	Leu	Met 230	Pro	Phe	Ser	Pro	Tyr 235	Glu	Pro	Leu	Asn	Phe 240	His	Ala	
_		_					_				_	_	-	gcc Ala	_	826
		cac					gcc					cca		gaa Glu		874
														cgc Arg		922
														tgc Cys 305		970
		_			_								_	gct Ala	_	1018
_					_	_			_	_	_			ttg Leu		1066
					_		_			_		_	_	ctc Leu		1114
			_	_		_	_			_				gtg Val		1162
	_	_			_			_	_	_			_	cgg Arg 385	_	1210
														ggt Gly		1258
			_		_			_		_				gtg Val	_	1306
_		_	_	_			_					_		acc Thr		1354
				_	_	_	tac Tyr	_		_				gag Glu	tga	1402
gato	gtgga	atg 1	ttgci	ttttg	gc a	cctt	acgg	g gg:	catci	ttga	gtc	cagc	tcc (cccci	aagatg	1462

agetgeagee	ececagagag	agetetgeae	gtcaccaagt	aaccaggccc	cagcerceag	1522
gcccccaact	ccgcccagcc	tctccccgct	ctggatcctg	cactctaaca	ctcgactctg	1582
ctgctcatgg	gaagaacaga	attgctcctg	catgcaacta	attcaataaa	actgtcttgt	1642
gagctg						1648
<210> 4 <211> 18 <212> DNA <213> Arti:	ficial Seque	ence				
<220> <223> PCR	Primer					
<400> 4 tccgtacgag	cccctgaa					18
<210> 5 <211> 23 <212> DNA <213> Arti	ficial Seque	ence				
<220> <223> PCR :	Primer					
<400> 5 tgagcctcgt	gtatcatctc	aag	,			23
<210> 6 <211> 21 <212> DNA <213> Arti	ficial Sequ	ence				
<220> <223> PCR	Probe			;		
<400> 6 tccacgccat	gttccagccc	t				21
<210> 7 <211> 21 <212> DNA <213> Arti	ficial Sequ	ence				
<220> <223> PCR	Primer					
<400> 7	ggtcgtattg	g				21

<210> 8						
<211> 26						
 <212> DNA	erandan nadi se s					
<213> Art1	ficial Seque	ence				
<220>						
<223> PCR 1	Primer					
<400> 8	+ a a > a + + + > a	an an art				26
ggcaacaaca	tccactttac	cagagu				26
<210> 9						
<211> 21						
<212> DNA	ficial Seque	ance				
(213) ALCI.	riciai beque	51106				
<220>						
<223> PCR 1	Probe					
-100- 0						
<400> 9	ccagggctgc	+				21
ogcooggood	ccagggccgc	ū				
<210> 10						
<211> 8133						
<212> DNA <213> Homo	saniens					
12237 1101110	baptons					
<400> 10		~b~~~~				٥.
gecatgttge	ccaggctggt	ctcaaactcc	taageteaag	taateeteet	accttggcct	60
cccaaattgt	tgggattata	gatgtgtgcc	actatgccca	gccaatgtaa	gattttgtag	120
tatattagtg	ttgctcctgt	cctctgctgc	agggcttttt	tgattgggac	tcagtgaatt	180
actccaatcc	ctgaagtcac	atcagttggc	ccttagccga	acaaaaataa	atatcattoo	240
geceedacee	cegaageeae	accageegge	ccccagccga	2093939033	acaccaccgg	210
tggccaaaga	tgacagtgaa	tgaacctgaa	atgttgggcc	ttgtgacttt	tgggcctcca	300
						2.52
ggtgtctcaa	aactgtcccc	catggaggga	gataaaagga	aagagcatgg	acctgacaga	360
tagaatacta	ggggctggtc	ccaqctqqqc	tgttggtcac	ttactatata	actottacao	420
ccatgggcag	ggcctggcct	ggctcaccag	ggggtgggag	gccaggaggc	cgtggccttg	480
			- h - h	.		E40
gtgagettet	cctaactgtg	cccatgotgg	ctgteccage	ctgaggagtt	cetgaaceag	540
agctcgccct	tctacttctg	gatgaatggt	gaccgcatcg	actccctgct	ggagaacgac	600
cggcagcaga	cgcacatgct	ggatgtcatg	caggaccact	tcagccgcgc	gtccagcatc	660
atagacgage	tcttccagga	caggttette	acccgggaage	cccaggatac	ctaccactac	720
						0

780 etgeeettea geetgeeeca eeggaggeet caettettet tteeeaagte eegeategte cgcagcttga tgcccttctc tccgtacgag cccctgaact tccacgccat gttccagccc 840 tteettgaga tgatacaega ggeteageag gecatggaca tecaetteca cageeeggee 900 960 ttccagcacc cgccaacaga attcatacga ggtgagaagg ggtggaagct catggccttt tgagcaactc gttagatgct gagaaccatg ccgagggctc agcgggtgtc atctcgattt 1020 ttctccagca atatcacaag ggtgatatta tccttattta aagaggaaaa aaactgagct 1080 gggcatggtg gctcatgcct gtgatgccag cactttgaga ggccaaggcg ggaggatcat ttgaggccag gagtttgaga ccagcctggc caagatagtg agaccctgtc tctacaaaaa taaaaactta aaaaattagc cgggtgtggt ggtgcacacc tgtagtctca gctactcggg 1260 aggetgagge aagagagtea eetgageetg gaagttggag getgeagtga getatgattg 1320 caccattgca ttccagcctg ggcaacagag tgagaccctg tctctaaatt aaaaaataaa 1380 taaaaataac aataggaatc agtggagtcc atctctgcat ggctggatga ctgactcttc 1440 ttccctcgtg tgtccccaga aggcgacgat gaccggactg tgtgccggga gatccgccac 1500 aactccacgg gctgcctgcg gatgaaggac cagtgtgaca agtgccggga gatcttgtct 1560 gtgggtgagt cggggtccag accacaagcc gtccccctg atcccttgtg tcctggggtc 1620 actggggcct cactggtgct gcctttatgg agtcagacag ataagcgttt ggattccagc 1680 tetgeageet ttgagetgtg teeeggggea ggteetgage cteatgeage tteggtteet 1740 1800 catcttagaa tgagatgatg atgcgaggct gtccctgaag tcggtgagat gtcgttagag atgcaaaagt gccctccacc tggtcggccc catgttgaaa aaagcttgtt gaaaaaagtc 1860 atccccctgg gactccccgg tgattctgtt cccaagcgcc aagcagtagg catcttcatt 1920 1980 ttcctctgca gattatgaca ttgcagacag tatgtgtttt gtttaacaaa actgaccaga ggccaggcac tgttctaaac actcgacata catttcctca tttcctcaga atgaccctct 2040 gaggaaactg agccacagaa aggttaataa cttatccaag attgaccccg acatgggcga 2100 gctgggcttc aatcctaggg cgctgtgttc tctcctgggg cccctcgcag cctctgccac 2160 agaagtcacg ggtctcagta cctgggcatc caagcaatag tccctttggt cggttggttg 2220 gtcccctagg caaagggaat atttcccttt aactgtcccc ctccgtttca ccagctctgg 2280 ttatgggtta acttetttee acttagagat aacagetgtg acagtatttg gaetagttee 2340 tggtacacag cagttcatac tcacaaagag ttaattgttt ccccttgttc aacagcttat 2400

cgatctggtg gctttgctct tacttaatgc ttagtttgag tttgccatgg caggccgcca 2460 gggtctagtt aaacattcct agcctcactc ctataatttt agaagccact gcaaaataaa 2520 cagttgtgct: ttaacaggct: gaagtataag: ttgctgtaga: tgagtgcaca: accaggcctt -- 2580 ggggcttttt ctataaaaaa tatcatagag tggcatcaat tacatggtac ctcaccacaa 2640 gaaagtcatg ttagggtctg agaaaagatg tcagatgcct gtgcccagat tggacctctt 2700 atagctgatt tttactctgt tgcccaggct gggtcaggtc tggcccaatc ttaacagtca 2760 ttgattacag ttgagagtgc agccagcgcc agtcttatca gtcattgatt atagctggcg 2820 tacaqtgqct ctatctcggc tcactgcgac ctccgcctcc tgggttcaag tgattctcct 2880 gcctcagcct cccaagtagc tgggagtgca ggtgtgcacc accacaccca gctaattttt gtatttttag tagagacagc atttcactat gttggccagg ctggtcttga actcctgacc 3000 tcaagtaatc tccccgcctc ggcctcccaa agttctggga ttacaggtgt gagccactgt 3060 gcctgacctg agatagattc ttagagaatt attggtaaga ataattctct aagctgagct 3120 aaatagtota cactgaagag gactgootac tgttatttaa ggtgottgoa accatataag 3180 catgtactgc ctgggaactc tagatgagga tttctcaatt tcagcgctgt tgattttttt 3240 tttttttttt gagacagggt ctctctctat cacccagcct ggagtgcagt ggcaccatta 3300 cageteactg cageetagae etettggget gaagteatee teetgeetea geeteetgag 3360 taacagacta caggtgtgct ccaccatgct tggctaattt ttttattttt agtagagatg 3420 gggtcttgct acattgccca agctggtctc taactcctgg gctcaagtga tcctcctacc 3480 tcagcctccc agagtgctgg gattacaggt gtgagcagtg ctgacatttt ggaccaggtc 3540 attetttgtc gttgggggct gtcctgagca gttcagggtg tttggcagca ttcctggcct 3600 ctgcccacta gaggtcagca gctcccttcc ctttgttgtg acaaccagct tcagaacttg 3660 ctaaatctcc ctgggtgaca gcgtccacag tagagaacct ctattctaga ctaagcctca 3720 gctcttaagg atttttctta ttttattatt atttttttaa gacagggtct cgctctatca 3780 cccaggctgg agcgtagtgg cgcaatcttt gctcactgca acctctgctt cctgggttca 3840 agogatttet cetgeeceag ceteetgagt agetgggatt acaggegtge acegeeaege 3900 ctggctaatt tttatatttt tagtagagac agggtttcac catattggcc aggctggtct 3960 caaactettg accteaagtg atcageetge etcageetee caaagtgetg ggattacagg tgtgagccag cacgcctggc tagtttttct tatttttaaa tttttttg gtaaaataat 4080

qatgtttatt tattacatat ttattttcaa actggcatct tgttaqtaat tctqtttctt 4140 tccccaccta acattttgtt tactataaat gatttcagtc atcatcctaa agcatatgca aaateteeet teeeetgaet eaegtttgat gtaeetgeet etggatattt ttgaaataee 4260 ttagggggag aaaaacagta gttttaagag ctagtggaca gtttccaggt cttaatgaat 4320 ctgacaacct gcagcccagg gccaagagga atgaattctc ttttccctgc tctcttgatg 4380 aactcactga ccagccatgg gcggcaggtg ggcaggcaag gacccctggc caccaggtgc 4440 cagtgcatca gctgcatgaa ctcctggcac cagaactgcc acctctacag acatgctcaa 4500 aagacaagtt tggaccgggt gcattggctc acacctgtaa tcccagcacc ttgagaggcc gaggtqqqtq gacccctgag gtcaggagtt tgagaccagc ctagccaaca tggtgaaacc 4620 ctgtctctcc taaaaataca aaaaaatcac ccgggggtgg tggcaggcac ctgtaatccc 4680 4740 aactactetg gaggetgagg caggagaatt gettgaacce gggaggtgga ggttgeagtg agctgagete gegecattge actecageet gggaaacaag agegaaatte tgteteaaaa 4800 aaaagacaag cttggaggat tgtccagaac cacagatcca gggtaggaaa agcccaagct 4860 taggagetga agaccetggt teaatceegg geceagagat catttattet atggetttag 4920 gtaagctatt tattgatact tctgtgggcc tcagtttcat tattggtaaa aattatttca 4980 ttattggtaa aattaggact taagtootaa toottaagto agaacagato caattottag 5040 agaaaaagga tatccagaga gaactttctg cggtgtctgg gacgcaggca gtgccacacg 5100 aatggcagct gtgagtaata ttcctcctct ctggaaatga ttcccgggag gactagggca 5160 5220 acgagagcca ctccaggtct gagaacatgg agaacttgag atcagtgctt ttggaagtgt ggtcaacaca gtttgtcacc aaagagataa gggtctggca cccaaagata aatgaatgat 5280 gttacgaagc acactgttta ggtcagttgg cgtatttttc cagagcaagg cttctcaggc 5340 tgggegtggt ggeteacace agtaateeca geactttttg ggeagatggg ttgageecag 5400 gagttcgaga ccagcctgga caacacagag aaaccccgtg tctacaaaaa atacaaaaat 5460 tagctgggca tggtagcatg tgcctatagt cccagctact caggaggctg aggttggagg 5520 5580 acageetgag eetgggaagt caaggetgea gtgageegag ateteaceae tgtatteeag cctaggcaac agagcaaaac tctgtctcaa aaaaacaaaa acaaaaacaa aaaacccaaa 5640 agactttctg gatgacggaa gcagtgtcta gattcacatt ctgaggcaaa acctttattt 5700 tgtcgtggac aattccagtt tgtggccctt cccttaggga agcactgctt ttgttcccgc

tgcatgtgct aacttccatt cattcatggt tctatccett tgtagccttc ccttcacact 5820 teteaettge gtttetteea tetetgggea gaetgtteea ceaacaacce eteccagget 5880 aagctgegge gggagetega egaateeete eaggtegetg agaggttgae eaggaaatae 5940 aacgagetge taaagteeta eeagtggaag atgeteaaca eeteeteett getggageag 6000 ctgaacgage agtttaactg ggtgtcccgg ctggcaaacc tcacgcaagg cgaagaccag 6060 tactatetge gggteaceae ggtgagetgt gteeeggeea catgetgtgg etegggagee 6120 gagctgtgat cgggagcagg ggcatgtgtg cttttgactg agcatttatc acacggcaga 6180 aaatagaaaa ctttaggcgc ccctgttgcc ttgaagcctc atcacccact cagggaaaat 6240 ataaccctgc tttacaaagg agcaaagtaa gagaggttcc acagcttggc caaggtgtga 6300 tagetgaeag atgaettgga egggtatttg aacetgaetg eetggetgee aageetgtat 6360 tttgttgttg ttgtttttgt tttggtgcac aaatctgtga ataaaccaga agcctctgtt 6420 cttttctcaa agctacaagg ctgccctctg gcatgtaaaa tggcttatga attagtacat 6480 cactetetge cagtgataaa aacttetete taggecagae atggtggete atgeetgtaa 6540 tcccagcact ttgggaggca gaggcaagag gattgcttga ggccaggaat ttgagaccag 6600 cctgggcaac acagcaagat tccctctcta caaaaaatac aaaaatcagt caggtgtggt 6660 ggcacacact tgtagtccca gctattcagg aggctgaggt gggaggattg cctgagccct 6720 gaagtggagg ctgcagtgag ctgtgatcac gccactgcac tccagcctgg gtgacagagt 6780 gagactetgt etettaaaaa atatatatat ataaaataat aaaataaagt taaaaaatea 6840 aataaaactt atttctagta ctgggaactc ttcttttct tttctttctt ccctccaggc 6900 cctctggatt ccttttctac cctactctga ccaagggctg cctaaagcaa atgtttggaa 6960 accactttta ttctttgggg tgctccctgg ctggtcattt gcagatgaca tttgccccaa 7020 cacatgagtg tctgtgaacc aggtccgttc tgtccactga gctgtactta cgtctagatg 7080 tataagaagc atggggtcag ctctctaggt tccttggagg agcaggagga cttccttatc 7140 agaagcctga cttctgttgc agagcgcatg cattttgacc acagtgtttc agctcttccc 7200 ttttctcttg ttccatttag gtggcttccc acacttctga ctcggacgtt ccttccggtg 7260 tcactgaggt ggtcgtgaag ctctttgact ctgatcccat cactgtgacg gtccctgtag 7320 aagtotocag gaagaaccot aaatttatgg agaccgtggc ggagaaagcg ctgcaggaat 7380 accgcaaaaa gcaccggtaa gcaggcgggc ctttcctgcg gcctgcaggg cccagtgagt 7440

ctctgggagc	cacaaaaaa	caaacaaagt	gcagactcta	tagcctggtg	ggaacgactc	7500
cgcccggagc	cagagcccaa	gaacaaagcc	aggaagttac	gggggaattt	tatttttcct	7560
ttggaggatg	ttttactttg	gaggataact	gttttttatt	tcagggagga	gtgagatgtg	7620
gatgttgctt	ttgcacctac	gggggcatct	gagtccagct	cccccaaga	tgagctgcag	7680
cccccagag	agagetetge	acgtcaccaa	gtaaaccagg	ccccagcctc	caggccccca	7740
actccgccca	gcctctcccc	gctctggatc	ctgcactcta	acactcgact	ctgctgctca	7800
tgggaagaac	agaattgctc	ctgcatgcaa	ctaattcaat	aaaactgtct	tgtgagctga	7860
tegettggag	ggtcctcttt	ttatgttgag	ttgctgcttc	ccggcatgcc	ttcattttgc	7920
tatggggggc	aggcaggggg	gatggaaaat	aagtagaaac	aaaaaagcag	tggctaagat	7980
ggtataggga	ctgtcatacc	agtgaagaat	aaaagggtga	agaataaaag	ggatatgatg	8040
acaaggttga	tccacttcaa	gaattgcttg	ctttcaggaa	gagagatgtg	tttcaacaag	8100
ccaactaaaa	tatattgctg	caaatggaag	ctt			8133

<210> 11 <211> 940

<212> DNA

<213> Homo sapiens

<400> 11 60 aagettgaac tggagcaagg gtaggcactt gcatgctggg tggccagcct atgggaaggc tcgccctggg gcagagggcc tggcacccag cagctctttg agtgcatgag cctgtggtct 120 180 ctgtgtgctc agccagcctt gtgtcttcct gtaggatgcc ctaaatgaga ccagggaatc agagacaaag ctgaaggagc tcccaggagt gtgcaatgag accatgatgg ccctctggga 240 agagtgtaag coctgootga aacagacotg catgaagtto tacgcacgog totgcagaag 300 tggctcaggc ctggttggcc gccaggtgaa aaggggacac atgagtggcc aaggctctga 360 gtggggaagg aggggagcct agtgaaatat gcttcattcc gcatgccaga tgcaattgat 420 tagcattggc tggcttgccc agagtgccat gctccattgg taatgtctgg catgagtaga 480 gagagtggag tcatcaaaag gatgtaggcc aggtatctgc cttctcttag aaaactcatg 540 cagcagtgct tagctggatg acataataaa ctgcttcgtg ggatgcagag ccctgtgtca 600 cttatgtgga aggatttaag aattttttt tttttttgag acagggtctc actctgtcac 660 ccaggctgga gtacagtgat gtgatcatgt ttcactgcag cttcgacctc ctgggttcag 720

780

gtgatcctcc cacctcaqcc tcccaaqtaq ctqqqactac aggcacgtac caccacaccc

agctaatttt tgtatttttt ttttgtaaac atggggtttg gccatgttgc ccaggctggt B40 ctcaaactcc taagctcaag taatcctcct accttggcct cccaaattgt tgggattata 900 gatgtgtgcc actagtccca gccaatgtaa gattttgtag 940 <210> 12 <211> 7610 <212> DNA <213> Homo sapiens <220> <221> unsure <222> 5461 <223> unknown <220> <221> unsure <222> 5462 <223> unknown <400> 12 gacctgcagg tcaacggatc cattcccgat tcctcatcgt ccagatggaa gaaactgagg 60 cccaagggca aagtgattag tccgaggtca cccagtgtct aggggcacac ctaggactgt 120 aatcagactt tcatggacct ggtctgggtt ctcccactta gtcatgggcc ttgaagattc 180 cccgaggctg cctcctgaaa aggactgggg tctagtggcc cctggacgtt gggcaagcaa 240 gggactgggc ctccatgttg tgcctccata gtcctgatcc tgaactggaa aactcagccc 300 ctgaccacgc agctctcctt taagcccctt tgtttcacat ggttttcaaa gtctgccacc 360 cacagtgggg ctgcctgtac ccgccctgtc cacccattgc cccagctgtc agccccttga 420 cttctctcct ggggcttaaa catccctggc tccaaaatgg gcagctcact ttcttcccca 480 agaagtaget geaceteeag ggtteetaga tttgeeeete ettgeeaggg ggaggggtgg 540 ctgcgacagg agattctccc tgctctcagc agaaggaact ccagcagttg gagaccagca 600 aacccctctg gacacagatc tgatttccta actgggaagg ctcagggcaa aataaaaatt 660 caggtccact ggttcaaaaa ctatgaagaa tttcaagacc gtcacagtag cccattaaac 720 caaacgtgga totqcaaggg toccacagec atgaagccca coetgettgg ttgggttoca 780 aaaagatggg gacagtgatt gcttaagctc tgtggatcaa ggaccccgga gaggccttct 840 ggetetecae atatetgete tgateactee taaacacaat tetgttteet ecaggeetgg 900

cgggtcagtc	cagggacccc	catcagtgtg	atgtttccag	gagtaggcgt	ttcaatactt	960
cctgtgctct	cttctccagc	acaaggcccc	tctccatccc	accctcatta	tgtctgactc	1020
tttactattt	aaatgggtca	agagaagtgg	cgcttgtgta	atgtgaaggt	taaggtcagt	1080
agggccaggg	aactgtgaga	ttgtgtcttg	gactgggaca	gacagccggg	ctaaccgcgt	1140
gagagggctc	ccagatggca	cgcgagttca	ggctcttccc	tactggaagc	gccagcgccg	1200
cacctcaggg	tctctcctgg	agccagcaca	gctattcgtg	gtgatgatgc	gccccccgc	1260
gccccagccc	ggtgctgcac	cggcccccac	ctcccggctt	ccagaaagct	ccccttgctt	1320
teegeggeat	tctttgggcg	tgagtcatgc	aggtttgcag	ccagccccaa	aggtgtgtgc	1380
gcgaacggag	cgctataaat	acggcgcctc	ccagtgccca	caacgcggcg	tcgccaggag	1440
cagcagcatg	ggcacagggt	ccgtgaccgg	tgagatgtcc	ccgtcttccc	tacccttgag	1500
cagagccaca	ccaggacgga	tgggcgggca [.]	ggggatggca	gccaggcaga	gagggatgac	1560
acagctegea	gtcacaaccc	ctgcgctttc	gacggagccc	aggaagccag	ggaggggagg	1620
tgccggagcc	ccatcaccag	gcagctgagc	caggggccgc	gcaaccgccg	cctgatgagc	1680
acgagcttca	cgcaaccaca	attctgtggt	ggggggtaa	atagaacaga	tataatgatc	1740
atcctttcgc	aaagatgggg	aaactgagac	ctggagacct	gccgcgttgg	cagacccagg	1800
ctagcaggtg	acagagctgg	cctgcaccga	getecttect	gcagcatatc	ctctgcgaag	1860
atgcggatct	ctcagttgtg	gctttcggct	tgcatgcatg	agtcatctag	ttttcttcta	1920
aattctctag	ctctctggac	actgttgcct	gtaagtatga	ggctgcggat	ttcagtatat	1980
ctgcaaccac	cgaaatccga	ctttttctgc	ctcctaatgc	atctgaggtg	catcagagaa	2040
aagtcacaca	agatccacca	ggcctcagac	ctctgattcc	acagteteat	tttacagatg	2100
ataatctgag	geetggagag	gtttaggact	ggtgccaaca	ctaaacagca	aataagtatc	2160
agaattggga	ttcgagccaa	agcctcttga	ccttccagaa	tttctggacc	tagttaaaaa	2220
aaatatgatt	tttattatta	tttttaaac	ggagaggtta	ggaatttaaa	ggaaagtaca	2280
gatactatat	aaaaaaagat	gcccatgaaa	atgttaagtt	ataataatag	tggagcattg	2340
ggcacaactg	aaatggccaa	tcttgtgaga	atggtaaaat	aaacttaggt	ccgtgagtaa	2400
gtggagtatt	acatagccat	aaaagtatgc	ccttaaagaa	tatttgaaga	tggtgaatgt	2460
gaagaatctt	gtataaactg	catggaagac	agaaggaaat	ataccacagt	gctaaccttt	2520
gcctctgggt	gatatgaatt	accggtgatt	atttttctta	ttttcctttt	ggtttagttt	2580

tctccatttg aagaagcaga taggagccgg ggctttggga ttgaaaccct caccatctgt gtgccctctt cactgtcttc ccatcctccc cacggctccc tgttcacagt cattgatttt etttetttet tttetettt ttttttttt teetgagaee aagteteaet etgttgeeca 2760 ggctggagta gagtagcgcc atctcggctc actgcaacct ccgccatccg ggttcaagca 2820 2880 gttctcatgc ctcagcctct gagtagctgg gactacagcc gcatgctgct acatccggct aatttttgta tttttagtag agacatggtt tcaccacctt ggccaggctg gtctcgaact 2940 cctgatctca agtaatccag cctgtcttgg cctcccaaag tgctggggtg acaggtgtga 3000 atcaatgcgt ccctgccagg tcattgattt tcttaagcct ctagccctgc cctgcttgga 3060 aacgttttgg gaagctgete agttcaaagt teecaggagg gtgtgeetgg aggggagttg 3120 ctcccaaagt ctgcctgctc ccccgcccc ccctgcccc cacccccgc catcttctcc 3180 3240 tecteetett eeeetgagea geeeetttgt eeaeagaace ggeettttet ggtagaagga gcaaggccaa gtggtttaag ccttcttagg gagaatgagg ctgtgtggta gtgctgggga 3300 ctcgagggcc ttgcgttggc atggctcttc cacccagggc agctggcagc caggctccca 3360 ggaggcagag gagatgaggg gggaggtgag tccgagcaaa ggaaaggagg tcggctgtgc 3420 agtcacggtt ctagaacatt cattggatca gcagcatcca tatcacctgc agactggctg 3480 gaaaagcagt ctcagaacca acattataac cagccctgca gtgattcata agtactttaa 3540 aaagtggtca atcatttcag caaagcagag ccacacagtc cgggggacca caggtggcct 3600 ctgtgtgctt gtctcggttt tcctgcccct ctccagacat gttgattaga cactgccaat 3660 gcccagcctc agacctcagt ctaatttgga agtagtcaga atttactatg attacataag 3720 accetegtgt ttacagaaca catteccete tetgaggtet ggattagate cattttacag 3780 atgaagaaac tgaggctcag atatttaagt gacttggaat caaggaaaga atactggaca 3840 3900 tggggctggg agggctgggc tctcatccca gggttaccat gagcatgctg tggactctag ggagtccatg ccctctctgg cgttcagctc accgctaggt agagaggttg ggtgagagaa 3960 cgacctcctt cccaggtctg agetggatgg ttcaccaggg accccaggct ccctggacga 4020 gactotgtgc ccgctgctga gtctggaatt cctttcctgt atcttgcctt tgcgtgcccc attetteatg geccageace etgetetetg gteagaacet agttetgaat gggtttttee agaagttgtt gettteaggg geeeetggea gagaggtgtt tetggetgge tttgtetete tggcatgaca aaggctctgt teetgetgga ggcattteag ggeteagtgg geagetgggg 4260

cagacgctga gaccacagcc ttcctggtga gcccggtctc cgccccctac cccatctctg 4320 ggaaggeget gaccecatet etteteccae getgetecet ggetetttge geetgattae 4380 ttctcatgag aggcactcct tgttaatgtg ctactgagtg tccagatggg cctgctgggc 4440 tgagcgggct ttggatgtga accatttcag gaaggggaac ccatcgtcct gttggttctg 4500 tgatggcaaa tgggtgagct cagataacga gttcttggga ggggcatggt gggggtggag 4560 tgcaggggga ggggtttctg ttttattgac aacagcctca gcttctggga aagggtccat 4620 4680 tgtgtaagac cggggctatg gctgtgcccc gtggctcagg gcagccagec agtggtggca ggaacactgg cagggcagcc tcgtgtcggc ttagagggga tgggcagtgt ggagggcctg 4740 gcagagcaag aggactcatc cttccaaagg gactttctct gggaagcctg ctcctcgggc 4800 cactgcgaac cetetetact etecgaagga attgteette etggetteea etactteeac 4860 4920 ccctgaatgc acaggcagcc cggcccaagt ctcccactag gatgcagatg gattcggtgt 4980 gaagggetgg etgetgttge etcegegtet tgaaagteaa gtteaggtgg tgetgagaet ccctgggggc tgcagcgctg tggtgaatgg ggagcgtctg ctggggtgaa ggtttaggtg 5040 cacattgcag aggacgtggc tggtctctgg gatgcagtcc ctctgtggag gtggcatggg 5100 gagggacgga tgcatgacct aagggtggta ttttcagtgt ctgacatgat cgataccact 5160 ctggacaagg aggccaggat gcagaaagcc tgtgtgcctc gctgattgtc ggggaggatg 5220 tggcttggac aagagcctgg ttcctccgat gccagggttc ttgtttcttc cactcaacat 5280 tgctgtcctg cagtccctcc ctcctgcac ctcctgcctt cgctttcatt cgaggtgtcc 5340 atggcaagtc tggtcatttc ccccatttc ctcaggaata aaagttgcag cagtgcctgc 5400 tgtggggaca gctgagggca gtgaggctgg ggagctgctg cagggcggag tgggcggac 5460 nnagcagget gtctagetgt teccatgatg gteteetgtt etetgcagag gegtgeaaag 5520 actecagaat tggaggeatg atgaagaete tgetgetgtt tgtggggetg etgetgaeet 5580 gggagagtgg gcaggtcctg ggggaccaga cggtctcaga caatgagctc cagggtgagt 5640 5700 agaccaagca tgatgttcct ctggccacag ggtgatgagg tcagagggca gggtagctaa ttctgctcag tgcctctcta tcaggcccca gtgttacaga ccgtttttat cttgtgcact 5760 gggtctgggt gcctgtgtct gggcccactc tgagcctcag ctcccaggcc cctggttcag 5820 getetgegtg cateagactg ceggeatttg caggeattte ceaageactt teggetgttg 5880 catttcattc agotottocc otoccaggod cottagedda getoccaggo ctoctedada 5940

aagetgtgte tggaecaceg gagetettat ceeteteece tttggagtge eeagagetta 6000 tccctcctgt gagctgacgg tttctgcagg atcattgtta aaaacccaga tcagacatgg 6060 gtgtgagtet gtttcacctc ttctcagetg ggtgactttg ggccactatc ttgatctcat 6120 gacactcccc ccaccccca ttttattgag atataattaa caaataaaaa ttgtgtatat 6180 ttaaggtata tgacgtgatg ttttgaaatg cacatacatt gaaatgatga ccagttttta 6240 tggtgggacg gtgggaagac ttaaaatcta ctttcttagc aaatttccag ttatgatatg 6300 gtgttattaa ctataagcac cacctgtatg ttagacctcc agaacatact cctcctacct 6360 gatgaacact ttgacccttt atcatatcac acttcccatg tctccctctg cgaagtgggc 6420 acggcggggg gctggagcat tacgtaaact gcacatgaag tgtttggcgc agtgcttggc 6480 atgggataaa caccaqtgaa qtaqcactta qqtgacacaq tqtttcqctq catttgtcac 6540 cagtgctatc cttactcatt tactcatctt cttattcctg tcgcctggca ctgcattgga 6600 acaaagaaat acacatatct gtttaaactg aactctagaa agatttgtgt ccaaaataac 6660 aatattttat attttgatgc tgcaaacgtg acacttctgg gttttttttt tttccttgcc 6720 aagtttette tgcacccage teatteteca ggggcacatg gcagtggctg ggcataacte 6780 tgggtgtgcc ggctcccatg gtctgcattt ctaagcagta gggtgcagtc agcaaggagc 6840 ctgtgatggg agcctgtgcc agggcaaggc tggggcatgc tgctgcctgc tggcaggagt 6900 gggggtccca gccttgacag cccctgaact gaacgggcct ttctggcatc cagctcattc 6960 7020 cagggtcctg aggccacctc ttcctctcgc ctcattctgc ctcttgcact tctcttgcag 7080 aaatgtccaa tcagggaagt aagtacgtca ataaggaaat tcaaaatgct gtcaacgggg tgaaacagat aaagactoto atagaaaaaa caaacgaaga gogcaagaca otgotcagca 7140 acctagaaga agccaagaag aagaaagagg tcaggaggag ccgctaccgc ctccctgcct 7200 tgaccatece actggagggg agggaggggg teactgegeg gtgecetget ggttgecatg 7260 gtgaccegca gtecteccag getgtgteag etgatgetga ggetgeagtt aagaageagg 7320 gaaggttcat ttgcttctga aagcatcagg gagtgagatc ttggatctgg ttttgttatg 7380 agcctggccc agggctaatg ccagattcat ttcaatagat gtttctaagc cctgatcacg 7440 tgctagttcc aagcaggctc tgggtggggt ggcggcaggg gccagacagg cgtggcgtcc 7500 aaccttcagg aagcttctag gagttaggga acagttggat cttgaaggat gagtgggttc 7560 tttaagccag gtgggaaggg gattccaggt gggcgaatga ggggaagctt 7610

<210> 13 <211> 1651 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (199)...(1545) <400> 13 ctgcgaaccc tctctactct ccgaagggaa ttgtccttcc tggcttccac tacttccacc 60 cctgaatgca caggcagccc ggcccaagtc tcccactagg gatgcagatg gattcggtgt 120 gaagggctgg ctgctgttgc ctccggctct tgaaagtcaa gttcagaggc gtgcaaagac 180 tccagaattg gaggcatg atg aag act ctg ctg ctg ttt gtg ggg ctg ctg Met Lys Thr Leu Leu Leu Phe Val Gly Leu Leu ctg acc tgg gag agt ggg cag gtc ctg ggg gac cag acg gtc tca gac 279 Leu Thr Trp Glu Ser Gly Gln Val Leu Gly Asp Gln Thr Val Ser Asp aat gag ctc cag gaa atg tcc aat cag gga agt aag tac gtc aat aag 327 Asn Glu Leu Gln Glu Met Ser Asn Gln Gly Ser Lys Tyr Val Asn Lys 30 gaa att caa aat gct gtc aac ggg gtg aaa cag ata aag act ctc ata 375 Glu Ile Gln Asn Ala Val Asn Gly Val Lys Gln Ile Lys Thr Leu Ile 45 50 gaa aaa aca aac gaa gag cgc aag aca ctg ctc agc aac cta gaa gaa 423 Glu Lys Thr Asn Glu Glu Arg Lys Thr Leu Leu Ser Asn Leu Glu Glu 65 gcc aag aag aag aaa gag gat gcc cta aat gag acc agg gaa tca gag 471 Ala Lys Lys Lys Glu Asp Ala Leu Asn Glu Thr Arg Glu Ser Glu aca aag ctg aag gag ctc cca gga gtg tgc aat gag acc atg atg gcc 519 Thr Lys Leu Lys Glu Leu Pro Gly Val Cys Asn Glu Thr Met Met Ala ctc tgg gaa gag tgt aag ccc tgc ctg aaa cag acc tgc atg aag ttc 567 Leu Trp Glu Glu Cys Lys Pro Cys Leu Lys Gln Thr Cys Met Lys Phe tac gca cgc gtc tgc aga agt ggc tca ggc ctg gtt ggc cgc cag ctt 615 Tyr Ala Arq Val Cys Arg Ser Gly Ser Gly Leu Val Gly Arg Gln Leu 125 130 gag gag ttc ctg aac cag agc tcg ccc ttc tac ttc tgg atg aat ggt 663 Glu Glu Phe Leu Asn Gln Ser Ser Pro Phe Tyr Phe Trp Met Asn Gly 140 145 150

gac o	Arg	Ile	Asp	Ser 160	Leu	Leu	Glu	Asn	Asp 165	Arg	Gln	Gln	Thr	His 170	Met	711
ctg c																759
gag d Glu I			Gln													807
cac t His 7		_			_	_										855
ccc a Pro I 220																903
ccc o	_				_	_		_					_			951
gag g Glu <i>l</i>	_	_	_	-	_	_					_	_	_		_	999
cac o	_			_			_	_		_	_	_				1047
tgc d Cys 1				_				_		_	_			_	_	1095
cag t Gln (300																1143
aac d Asn I			_	_		_					_	_			_	1191
gtc q Val 1																1239
cag t																1287
cag t Gln 1							-	-			_		_	_	_	1335

											cac His					1383
											aag Lys				tct Ser	1431
											tcc Ser					1479
											cag Gln					1527
			gag Glu		tga	gat	gtgga	atg 1	tgct	ttt	gc ad	ecta	eggg	g gca	atctgag	t 1585
cca	gete	ccc (ccaa	gatga	ag ct	gcag	gccc	c cca	agaga	agag	ctct	gca	egt o	cacca	aagtaa	1645
cca	ggc															1651
<213 <213 <213 <220)>	O NA rtif:			quen											
<22	3 > A1	ntise	ense	Olig	gonu	cleo	tide									
	0> 14 cttg		gcct	cggt	ca											20
<21 <21	0> 19 1> 20 2> DI 3> Ar	O NIA	icia:	l Sed	quen	ce										
<220 <220		ntis	ense	Oli	gonu	cleo	tide	,								
	0> 1! ctgga		cttt	gcac	gc											20
<21:	0> 10 1> 20 2> DI 3> Ar	O ATA	icia	l Sed	quen	ce										
<22	0>															

<223> Antise	ense Oligonucleotide	
<400> 16 gtcttcatca t	gcctccaat	20
<210> 17 <211> 20 <212> DNA <213> Artifi	icial Sequence	
<220> <223> Antise	ense Oligonucleotide	
<400> 17 tctcccaggt o	cagcagcagc	20
<210> 18 <211> 20 <212> DNA <213> Artifi	icial Sequence	
<220> <223> Antise	ense Oligonucleotide	
<400> 18 tctggtcccc o	caggacctgc	20
<210> 19 <211> 20 <212> DNA <213> Artifi	icial Sequence	
<220> <223> Antise	ense Oligonucleotide	
<400> 19 ggagctcatt g	gtctgagacc	20
<210> 20 <211> 20 <212> DNA <213> Artifi	icial Sequence	
<220> <223> Antise	ense Oligonucleotide	
<400> 20 acttacttcc (stgattggac	20
<210> 21 <211> 20		

<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	***************************************
<400> 21 aatttcctta ttgacgtact	20
<210> 22	
<211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	•
<400> 22 gtctttatct gtttcacccc	20
<210> 23 <211> 20 <212> DNA	
<213> Artificial Sequence <220>	
<223> Antisense Oligonucleotide	
<400> 23 gggcatcctc tttcttcttc	20
<210> 24 <211> 20	
<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 24 atttagggca tcctcttct	20
<210> 25 <211> 20	
<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 25	20

<210> 26 <211> 20		
<212> D		
2735 AI	tificial Sequence	
\Z_15/ III	corract poducing	
<220>		
	eticomo Olimonuslockido	
<223> AI	ntisense Oligonucleotide	
<400> 26		_
tgtctctg	gat tecetggtet 2	0
	· _	
<210> 23	•	
<211> 20		
<212> Di	AI	
<213> A1	rtificial Sequence	
<220>		
<223> Ar	ntisense Oligonucleotide	
<400> 27	7	
		0
<i>333</i> 3		
<210> 28	B	
<211> 20		
<211> 20		
	rtificial Sequence	
<213> A	refilerat sequence	
000		
<220>		
<223> AI	ntisense Oligonucleotide	
	_	
<400> 28		_
cccagag	ggc catcatggtc 2	0
<210> 29		
<211> 20		
<212> DI	NA A	
	rtificial Sequence	
	-	
<220>		
	ntisense Oligonucleotide	
<400> 29	9	
		0
		-
<210> 30	n	-
<211> 20		
<212> DI		
<213> A	rtificial Sequence	
<220s		

<223> Antisense Oligonucleotide	
<400> 30	•
tcaggcaggg cttacactct	20
 <210> 31	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 31	
gtgcgtagaa cttcatgcag	20
<210> 32 <211> 20	
<211> 20 <212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 32	
ggcggccaac caggcctgag	20
<210> 33	
<211> 20	
<212> DNA <213> Artificial Sequence	
(213) Attiticial bequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 33	2.0
tggcggccaa ccaggcctga	20
210. 24	
<210> 34 -211> 20	
<211> 20 <212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 34	
actcctcaag ctggcggcca	20
<210> 35	
<211> 20	

<212>		
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	35	
agtaga	aggg cgagctctgg	20
<210>	36	
<211>	20	
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	36	
ttcato	caga agtagaaggg	20
<210>	37	
<211>		
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	37	
cagcag	ggag tcgatgcggt	20
<210>	38	
<211>		
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	38	
gatga	ggtc gttctccagc	20
<210>	39	
<211>	20	
<212>		
<213>	Artificial Sequence	
<220>		
	Antisense Oligonucleotide	
<400>	39	
		20

<210>	40	
<211>	20	
<212>	DNA	
2135	Artificial Sequence	
~2107	TELLICIAL DEGACING	
000.		
<220>		
<223>	Antisense Oligonucleotide	
<400>	40	
gacato	cage atgtgcgtct	20
_		
<210>	4.1	
<211>		
<212>		
<213>	Artificial Sequence	
<220>		
	Antisense Oligonucleotide	
<400>	4.1	•
		20
grggro	ctgc atgacatcca	20
<210>	42	
<211>	20	
<212>	DNA	
	Artificial Sequence	
	•	
<220>		
	Antisense Oligonucleotide	
\2237	Ancisense Oligonacieotide	
400	40	
<400>	·	
aagtgg	tcct gcatgacatc	20
<210>	43	
<211>	20	
<212>		
	Artificial Sequence	
\Z13 /	Attitudat bequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	43	
ctqqaa	gagc tcgtctatga	20
<210>	44	
<211>		
<212>		
<213>	Artificial Sequence	
<220>		

<223> Antisense Oligonucleotide	
<400> 44	
 tgtcctggaa gagctcgtct	20
<210> 45	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 45	
gaacctgtcc tggaagagct	20
<210> 46	
<211> 20	
<212> DNA <213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 46	
ggaaagaaga agtgaggcct	20
<210> 47	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 47	
gggcatcaag ctgcggacga	20
<210> 48	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 48	
ctcaaggaag ggctggaaca	20
<210> 49 <211> 20	
SALLA AV	

<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 49 tctcaaggaa gggctggaac	20
<210> 50 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 50 tgtatcatct caaggaaggg	20
<210> 51 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 51 gctgtggaag tggatgtcca	20
<210> 52 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 52 attctgttgg cgggtgctgg	20
<210> 53 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 53 tatgaattet gttggegggt	20

<210> 54 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 54 ggatctcccg gcacacagtc	20
<210> 55 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 55 cggatctccc ggcacacagt	20
<210> 56 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 56 gtggagttgt ggcggatctc	20
<210> 57 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 57 gtccttcatc cgcaggcagc	20
<210> 58 <211> 20 <212> DNA <213> Artificial Sequence	
<220>	

20
20
20
20
•
20
_
20

<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 63 agctgctcca gcaaggagga	20
<210> 64 <211> 20	
<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 64 gctcgttcag ctgctccagc	20
<210> 65 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 65 ttgccagccg ggacacccag	20
<210> 66 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 66 cgcagatagt actggtcttc	20
<210> 67 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 67 accgtggtga cccgcagata	20

<210>	68	
<211>	20	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
	-	
<400>	68	
cgagto	agaa gtgtgggaag	20
<210>	69	
<211>	20	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	69	
gtgate	ggat cagagtcaaa	20
<210>	70	
<211>	20	
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>	70	
ggaga	cttct acagggaccg	20
<210>	71	
<211>		
<212>		
<213>	Artificial Sequence	
<220>		
<223>	Antisense Oligonucleotide	
<400>		
gccac	ggtct ccataaattt	20
		•
<210>		
<211>		
<212>		
<213>	Artificial Sequence	
<220>		

<223> Antisense Oligonucleotide	
<400> 72	
gcaaaagcaa catccacatc	20
J	
<210> 73	
<211> 20	
<212> DNA <213> Artificial Sequence	
(213) Artificial bequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 73	
	20
tagagtgcag gatccagagc	20
<210> 74	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 74	
attagttgca tgcaggagca	20
accagoogea ogeaggagea	-
<210> 75	
<211> 20	
<212> DNA <213> Artificial Sequence	
· · · · · · · · · · · · · · · · · · ·	
<220>	
<223> Antisense Oligonucleotide	
<400> 75	
agacagtttt attgaattag	20
2010. 76	
<210> 76	
<211> 20	
<212> DNA <213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 76	
cgagatagag ccactgtacg	20
<210> 77	

<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 77 tgccaccacc cccgggtgat	20
<210> 78 <211> 20 <212> DNA	
<213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 78 gttgttggtg gaacagtcca	20
<210> 79 <211> 20	
<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 79 tgcttaccgg tgctttttgc	20
<210> 80 <211> 20	
<212> DNA <213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 80 'acatctcact cctcccggtg	20
<210> 81	
<211> 20	
<212> DNA <213> Artificial Sequence	
<220> <223> Antisense Oligonucleotide	
<400> 81	
gaccetecaa gegateaget	20

C 0 9

<210> 82 <211> 20 <212> DNA	
<213> Artificial Sequence	•
<220> <223> Antisense Oligonucleotide	
<400> 82	
aaaaagagga ccctccaagc	20
	·
<210> 83	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 83	
tgtgtcccct tttcacctgg	20
<210> 84	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 84	
attaccaatg gagcatggca	20
<210> 85	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 85	
caacatggcc aaaccccatg	20
<210> 86	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	

<223> Antisense Oligonucleotide	
<400> 86 gcggcaggtc tccaggtctc	20
<210> 87	
<211> 20 <212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 87	
ttcccttcgg agagtagaga	20
<210> 88	
<211> 20 <212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
400 00	
<400> 88 tgcttgggaa atgcctgcaa	20
<210> 89	•
<211> 20	
<212> DNA <213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 89	2.0
agctggatgc cagaaaggcc	20
<210> 90	
<211> 20	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Antisense Oligonucleotide	
<400> 90	
tagaagtagt ggaagcagg	20

INTERNATIONAL SEARCH REPORT

International application No. PCT/US01/25255

	COLETA TION OF CUREOR LAND		
1	SSIFICATION OF SUBJECT MATTER	•	ſ
, , ,	:Please See Extra Sheet. :Please See Extra Sheet.		
	to International Patent Classification (IPC) or to bot	th national classification and IPC	{
		a national crassification and if C	
	LDS SEARCHED		
iviinimum a	ocumentation searched (classification system followe	• •	}
U.S. :	485/6, 91.1, 325, 375; 586/23.1, 23.2, 24.3, 24.31,	24.33, 24.5; 514/44	
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched			
Flectronic	data base consulted during the intermedianal according	name of data land at 1 at 1	
1	data base consulted during the international search (edline, caplus, lifesci, embase, uspatfull	name of data base and, where practicabl	e, search terms used)
C. DOC	CUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.
X	WO 00/49937 A2 (THE UNIVERSIT)	Y OF BRITISH COLUMBIA)	1, 2, 4, 5, 11
	31 August 2000 (31-08-00), see entire		
Y	, , , , , , , , , , , , , , , , , , , ,		3, 6-10, 12-20
			-,,
X	KANG ET AL. Antisense Oligonuc	eleotide of Clusterin mRNA	11
	Induces Apoptotic Cell Death and Pre		
Y	17D Sertoli Cells. Molecules and Cells		1-10, 12-20
	No. 2, pages 193-198, see entire docu		1 10, 12 20
	pages and and, see emile deep		
	•		
	L		
	her documents are listed in the continuation of Box	C. See patent family annex.	
-	ecial categories of cited documents:	"T" later document published after the inte date and not in conflict with the app	
	cument defining the general state of the art which is not considered be of particular relevance	the principle or theory underlying the	
"E" ear	rlier document published on or after the international filing date	"X" document of particular relevance; the	e claimed invention cannot be
	cument which may throw doubts on priority claim(s) or which is	when the document is taken alone	ten to maniac un magnita greb
	ed to establish the publication date of another citation or other scial reason (as specified)	"Y" document of particular relevance; th	e claimed invention cannot be
	cument referring to an oral disclosure, use, exhibition or other cans	considered to involve an inventive step with one or more other such docum obvious to a person skilled in the art	when the document is combined nents, such combination being
than the priority date claimed		"A" document member of the same patent family	
Date of the	actual completion of the international search	Date of mailing of the international se	arch report
23 OCTO	2S OCTOBER 2001 FEB 2007		
Name and mailing address of the ISA/US Authorized officer			
Box PCT	TARAKNA TACOHOCIEDE ACCO		
Washington, D.C. 20231			
Facsimile No. (703) 305-3230		Telephone No. (703) 306-0196	1

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US01/28235

C (Continua	C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.	
X · Y	MIYAKE ET AL. Antisense TRPM-2 Oligodeoxynucleotides Chemosensitize Human Androgen-independent PC-3 Prostate Cancer Cells Both in Vitro and in Vivo. Clinical Cancer Research. May 2000, Vol. 6, pages 1655-1663, see entire document.	1, 2, 4, 5, 11, 12, 14, 15 3, 6-10, 13, 16-20	
X Y	MIYAKE ET AL. Testosterone-repressed Prostate Message-2 Is an Antiapoptotic Gene Involved in Progression to Androgen Independence in Prostate Cancer. Cancer Research. 01 January 2000, Vol. 60, pages 170-176, see entire document.	11 1-10, 12-20	
X Y	MIYAKE ET AL. Acquisition of Chemoresistant Phenotype by Overexpression of the Antiapoptotic Gene Testosterone-repressed Prostate Message-2 in Prostate Cancer Xenograft Models.Cuacer Research. 01 May 2000, Vol. 60, pages 2547-2554, see entire document.	11 1-10, 12-20	
X - Y	ZWAIN ET AL Clusterin Protects Granulosa Cells from Apoptotic Cell Death during Follicular Atresia. Experimental Cell Research. March 2000, Vol. 257, pages 101-110, see entire document.	11 1-10, 12-20	
X Y	URBICH ET AL. Laminar Shear Stress Upregulates the Complement-Inhibitory Protein Clusterin A Novel Potent Defense Mechanism Against Complement-Induced Endothelial Cell Activation. November 1999, Vol.101, pages 352-355, see entire document.	11 1-10, 12-20	
X - Y	SENSIBAR ET AL Prevention of Cell Death Induced by Tumor Necrosis Factor alpha in LNCaP Cells by Overexpression of Sulfated Glycoprotein-2 (Clusterin). Cancer Research. 01 June 1995, Vol. 55, pages 2431-2437, see entire document.	11 1-10, 12-20	
Y	US 5,801,154 A (BARACCHINI ET AL.) 01 September 1998 (01-09-98), see especially columns 6-9.	1-20	
Y	MILNER ET AL. Selecting effective antisense reagents on combinatorial oligonucleotide arrays. Nature Biotechnology. June 1997, Vol. 15, pages 537-541, see entire document.	1-20	

INTERNATIONAL SEARCH REPORT

International application No. PCT/US01/28235

A. CLASSIFICATION OF SUBJECT MATTER: IPC (7):		
C07H 21/02, 21/04; A61K-+8/00; C12Q-1/68; C12P-19/84; C12N-1/85; -15/86		
A. CLASSIFICATION OF SUBJECT MATTER: US CL:		
+35/6, 91.1, 325, 375; 536/23.1, 23.2, 24.3, 24.31, 24.33, 24.5; 51.	1/++	
·		
	·	
,		
	·	
	·	

Form PCT/ISA/210 (extra sheet) (July 1998)*